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Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Run on:
                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                             Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                   Database :
                                                                                                                                                                                                                                                                                                              BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                January 6, 2003, 20:58:58; Search time 21 Seconds (without alignments) 375.493 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                US-09-674-235-1
1406
                                                                                                                                                                                                                                                                                                                                                              1 MAFEELLSQVGGLGRFQMLH......RKKISRKRHKNDCYTKVTKF 268
262574 segs, 29422922 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                       262574
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| 2254 | 16 19 20 21 22 | 10 11 11 13 14 | 7654822 | Result No. |
|--|--|--|--|---------------|
| 80.55 5.55 | 2. 88. 3. 8 | 146 104 104 100 100 20,5 | 458.5 436.5 369.5 169.5 152.5 | Score |
| 5.77 5.77 | 55 55 55 55 55 55 55 55 55 55 55 55 55 | | 32.6 31.0 26.1 12.1 12.1 10.8 | 당각 |
| 521 521 521 1308 | 581 494 492 492 523 325 | 2255555 222003 286644 | 55555555555555555555555555555555555555 | Length D |
| 2521 | 00440D40 | 30 CO 44 CO 44 44 | 4400000 | DB |
| -08-075-193- -08-564-090A T-US94-06698 -08-996-644- | US-08-989-386-7 US-09-031-392-5 US-09-299-549-5 US-09-610-417-5 US-08-355-844-3 US-08-355-844-3 US-09-295-16126-3 US-09-295-197-5 | -501-572- -040-444- -964-127- -496-692- -928-692- -339-972- -964-127- -496-692- | -09-330-245A -09-572-147- -08-647-397- -08-501-572- -09-040-444- -09-040-444- | ID |
| equence 2, equence 2, equence 2, equence 2, equence 2, | Appl Appl Appl Appl Appl Appl Appl Appl | Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 1, Appli Sequence 12, Appli Sequence 12, Appli Sequence 4, Appli Sequence 4, Appli | , | Description |

| 80.5 5.7 1308 3 US-09-352-552-2 80 5.7 1024 4 US-09-562-737-89 80 5.6 285 3 US-09-187-050-12 79 5.6 393 3 US-09-187-050-14 79 5.6 393 3 US-09-187-050-16 79 5.6 393 3 US-09-187-050-18 79 5.6 393 3 US-09-187-050-20 79 5.6 393 3 US-09-187-050-24 79 5.6 393 3 US-09-187-050-26 79 5.6 393 3 US-09-187-050-27 79 5.6 393 3 US-09-187-050-29 79 5.6 393 3 US-09-187-050-29 79 5.6 393 3 US-09-187-050-39 79 5.6 393 3 US-09-187-050-39 79 5.6 393 3 US-09-187-050-39 79 5.6 393 3 US-09-187-050-33 79 5.6 393 3 US-09-187-050-33 79 5.6 393 3 US-09-187-050-33 | 45 | 44 | 43 | 7.5 | ۱ د ا د | 41 | 40 | 39 | 38 | 37 | 36 | 35 | 34 | ω u | 32 | 31 | 30 | 29 | 28 |
|--|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|--|--|------------------|------------------|------------------|------------------------------|-------|
| | 79 | 79 | 79 | ! ; | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 79 | 80 | 80.5 |
| 3 US | 5.6 | 5.6 | ა. ი | | ת ת | Մ • თ | J. 6 | 5.6 | 5.6 | 5.6 | 5. 6 | 5.6 | 5.6 | 5. 6 | ა . | ა . რ | | | |
| SSD | 393 | 393 | 393 | 3 (| 202 | 393 | 393 | 393 | 393 | 393 | 393 | 393 | 393 | 393 | 393 | 393 | 285 | 1024 | 1308 |
| | ω | w | L |) د | w | ω | W | w | w | ω | W | w | W | ω | ω | W | | 4. | · W |
| | US-09-187-050-33 | US-09-187-050-32 | US-U9-IB/-U5U-31 | 10 00 101 000 01 | US-09-187-050-30 | US-09-187-050-29 | US-09-187-050-28 | US-09-187-050-27 | US-09-187-050-26 | US-09-187-050-24 | US-09-187-050-22 | US-09-187-050-20 | US-09-187-050-18 | US-09-187-050-16 | US-09-187-050-14 | US-09-187-050-2 | US-09-18/-050-12 | US-09-562-737-89 | -09 |
| | 30, | י י י |) L | <u>ب</u> | 30, | 24 | 200 | | 10 | 4 (| 4 . | 2 6 | 9 6 | 9 6 | , 1 | | ~ | 3 0 | |
| 00H000H10H00m0H-10A | TOGW | 1001 | 100 | Ann | Appl | Tddw | 1001 | 1001 | ייין ל | 1001 | 1001 | 100 | יייייייייייייייייייייייייייייייייייייי | יייייייייייייייייייייייייייייייייייייי | 777 | 1 7 7 | 7 | ביים ביים ביים ביים | Trada |

ALIGNMENTS

| Qy 193 KVARTNGIKNABETLNIEVVRSTMQEELDAAQTKTTVCDLFRNPSMRKRICILVFL 248 |
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| Qy 172VEŞARWLIITNKLDEGLKALR 192 |
| Db 170 LQTAVSGTCAAFAPNFPIYCAFRLLSGMALAGISLNCMTLNVEWMPIHTRACVGTLIGYV 229 |
| Qy 170 171 |
| Db 110 IYDNSTEPSTIVTEWDLVCSHRALRQLAQSLYMVGVLLGAMVFGYLADRLGRRKVLILNY 169 |
| QY 121 VYDQSYFPSTIVTKWDLVCDYQSLKSVVQFLLLTGWLVGGIIGGHVSDR 169 |
| Db 55 ANLSKNGGLEVWLPRDRQGQPESCLRFTSPQWGLPFLNGT-EANGTGATEPCTDGW 109 |
| Qy 61 GNETGIISEDALLRISIPLDSNLRPEKCRRFVHPQWQLLHLNGTIHSTSEADTEPCVDGW 120 |
| Db 1 MAFNDLLQQVGGVGRFQQIQVTLVVLPLLLMASHNTLQNFTAAIPTHHCRPPAD 54 |
| Qy 1 MAFBELLSQVGGLGREQMLHLVFILPSLMLLIPHILLENFAAAIPGHRCWVHMLDNNTGS 60 |
| Query Match 32.6%; Score 458.5; DB 4; Length 550; Best Local Similarity 29.2%; Pred. No. 4.5e-44; Matches 104; Conservative 51; Mismatches 82; Indels 119; Gaps |
| TYPE: PRT ORGANISM: Unknown FEATURE: OTHER INFORMATION: Description of Unknown Organism: This information OTHER INFORMATION: is not available. US-09-330-245A-2 |
| ; SOFTWARE: Patentin ver. 2.0 ; SOFTWARE: Patentin ver. 2.0 ; SOFTWARE: Patentin ver. 2.0 |
| SEQ ID NO |
| APPLICATION N |
| PPLICAT |
| CURRENT FILING DATE: 1999-06-10 |
| |
| US-09-330-245A-2; Sequence 2, Application US/09330245A; Patent No. 6432631 |
| RESULT 1 |

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Sequence 2, Application US/09572147

Sequence 3, APPLICANT: Lin Yue

APPLICANT: Lin Yue

APPLICANT: Lin Yue

TITLE OF INVENTION: ENCODING MURINE ORGANIC ANION TRANSPORTER 5 (mOATPS) AND

TITLE OF INVENTION: SCREENING MURINE ORGANIC ANION TRANSPORTER 5 (mOATPS)

TITLE OF INVENTION: SCREENING MURINE OF 572,147

CURRENT APPLICATION NUMBER: US/09/572,147

CURRENT PALLING DATE: 1999-06-19

NUMBER OF SEQ ID NOS: 2

SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     55 AN----LSKDGGLEAMLPLDKQGRPESCLRFPFPH-----NGT-EANGTGVTEPCLDGW 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 VYDQSYFPSTIVTKWDLYCDYQSLKSVVQFLLLTGMLYGGIIGGHVSDR-----169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   164 LQTAVSGTCAAYAPNYTVYCIFRLLSGMSLASIAINCMTLNMEWMPIHTRAYVGTLIGYV 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               224 YSLGOFLLAGIAYAVPHWRHLOLAVSVPFVAFIYSWFFIESARWYSSSGRLDLTLRALQ 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----VESARWLIITNKLDEGLKALR 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 MAFEELLSQVGGLGRFQMLHLVFILPSLMLLIPHILLENFAAAIPGHRCWVHMLDNNTGS 60
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; Sequence 2, Application US/08647397
; Patent No. 5972702;
; GENERAL INFORMATION:
APPLICANT: Brier, David R.
APPLICANT: Brier, David R.
TITLE OF INVENTION: OSTEOCLAST TRANSPORTER
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wolf, Greenfield & Sacks, P.C.
STREET: 600 Atlantic Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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OCGANISM: MUS MUSCULUS
US-09-572-147-2
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Matches 102; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 600 AUCITY: Boston STATE: MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE.
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
ZIP: 02210
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APPLICANT: Koepsell, Hermann
APPLICANT: Koepsell, Hermann
APPLICANT: Grudeman, Dirk
APPLICANT: Grudeman, Dirk
APPLICANT: Grudeman, Dirk
TITLE OF INVENTION: Transport protein Which Effects The
TITLE OF INVENTION: Transport Of Cationic Xenobiotics and\or Pharmaceuticals,
TITLE OF INVENTION: DNA Sequences Encoding It And Their Use.
CORRESPONDENCES:
ADDRESSE: Finnegan, Henderson, Farabow, Garrett & Dunner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51 VHMLDNNTGSGNETGILSEDALLRISIPLDSNLRPEKCRRFVHPQWQLLHLNGTIHSTSE 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                111 ADTEPCVDGWVYDQSYFPSTIVTKWDLVCDYQSLKSVVQFLLLTGMLVGGIIGGHVSDR- 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90 GATEPCLDGWIYNST--RDTIVTEMDLVCGSNKLKEMAQSVFWAGILVGGPVFGELSDRF 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     148 GRKPILTWSYLLLAASGSSAAFSPSLTVYMIFRFLCGCSISGISLSTIILNVEWVPTSTR 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ------WLVESARWLIITN 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             208 AISSTIIGYCYTIGOFILPGLAYAVPOWRWLQLSVSAAFFIFSLLSWWVPESIRWLVLSG 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | | | :| :| :| | | | | | | | :
1 MTFSEILDRVGSMGPFQYLHVTLIALPILGIANHNLLQIFTATTPDHHCRPPNASLEPW 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 V------NASLPNDTQ 89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26.1%; Score 367.5; DB 2; Length 537; 24.6%; Pred. No. 1.3e-33; Live 45; Mismatches 74; Indels 151;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1300 I Street, N.W., Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MAFBELLSQVGGLGRFQMLHLVFILPSLMLLIPHILLENFAAAIPGHRC--
                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/647,397
                                                                                                                                                     FILING DATE:

CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:

NAME: Gates, Edward R.

REGISTRATION NUMBER: 31,616
REPERENCE/DOCKET NUMBER: B0801/7048
TELECOMUNICATION INFORMATION:

TELEPHONE: 617-720-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08501572
Patent No. 6063623
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                     TELEFAX: 617-720-2441
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 537 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: protein
COMPUTER READABLE FORM:
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
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US-08-501-572-1
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TELEPHONE: (202)408-4000
TELEPAX: (202)408-4400
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: PATENT APPLICATION DATA:
APPLICATION NUMBER: US/08/501,572
APPLICATION NUMBER: US/08/501,572
                                                                                                                                                                                                                        sequence 1, Application US/09040444 Patent No. 6063766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 35,391
REFERENCE/DOCKET NUMBER: 02481.1453-00000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     109 SLVANRSQLPLGÞĆEHĠŴŸŹDTP--GŚSÍŸTEFNĹŸĆGDAWKVDLFQSCVNLGFFLĠSLV 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                103 GTIHSTSEADTEPCVDGWVYDQSYFPSTIVTKWDLVCDYQSLKSVVQFLLLTGMLVGGII 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          167 VĠYIADREGRKLCILVTTLVTSVSGVLTAVAPDYTSMLLERLLQGMVSKGSWVSGYTLIT 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 163 GGHVSDR------ 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50 --WVHMLDNN-----TGSGNETGILSEDALLRISIPLDSNLRPEKCRRFVHPQWQLLHLN 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 12.1%; Score 169.5; DB 3 Local Similarity 20.5%; Pred. No. 8.9e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                           APPLICANT: Koepsel, Hermann
APPLICANT: Grundeman, Dirk
APPLICANT: Gorboulev, Valantin
TITLE OF INVENTION: Transport protein Which Effects The
TITLE OF INVENTION: Transport Of Cationic Xenobiotics and or Pharmaceuticals,
TITLE OF INVENTION: DNA Sequences Encoding It And Their Use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           175 ARWLIITNKLDEGLKALRKVARTNG-IKNAEETLNIEVVRSTMQEELDAAQTKT-TVCDL 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               227 BEVGSGYRRTTAILYQMAFTVGLVGLAGVAYAIPDMRWLQLAVSLPTFLFLLYYMFVPES 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 CGWSQAEELNYTVPGLGPSDEASFLSQ--CMRYEV--DWNQSTLDC---VDP-----LS 108
                                                                                                                                                                                                                                                                                                                                               339
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                                                                                                                                                                                                                                                                                                                                                                                                                      287 PRWILSQKRTTRAVRIMEQIAQKNGKVPPAD-----LKMLCLEE-DASEKRSPSPADL 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 DDVLEQVGEFGWFQKQAFLLLCLISASLAPIYVGIVFLGFT---- PGHYCQNPGVAELSQR 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 BELLSQVGGLGREQ---MLHLVFILPSLMLLIPHILLENFADAIEGHRC------ 49
                                                    CORRESPONDENCE ADDRESS:
                                                                        NUMBER OF SEQUENCES:
STREET: 1300 I S
CITY: Washington
                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                 FRTPNLRKHTVILMYL 354
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     556 amino acids
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                E: Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P.
1300 I Street, N.W., Suite 700
                                                                                                                                                                                                 Koepsell, Hermann
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US-09-040-444-1
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TELEPAX: (202)408-4400
INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                            US-08-501-572-3
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NAME: O'COMNOY, Steven P
REGISTRATION NUMBER: 41,225
REFERENCE/DOCKET NUMBER: 2481.1453-01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                        Sequence 3, Application US/08501572
Patent No. 6063623
GENERAL INFORMATION:
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LENGTH: 556 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       109 SLVANRSQLPLGÞCEHGWVYDTP--GSSÍVTEFNLVCGDAWKVDLFQSCVNLGFFIGSLV 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       103 GTIHSTSEADTEPCVDGWYYDOSYFPSTIVTKWDLVCDYQSLKSVVQFLLLTGMLVGGII 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
            APPLICANT: Grundeman, Dirk
APPLICANT: Gorboulev, Valentin
TITLE OF INVENTION: Transport Of Cationic Xenobiotics and or Pharmaceuticals,
TITLE OF INVENTION: Transport Of Cationic Xenobiotics and Their Use.
TITLE OF INVENTION: DNA Sequences Encoding It And Their Use.
                                                                                                                                                                                                                                                                                                                                                        227 EFVGSGYRRTTAILYQMAFTVGLVGLAGVAYAIPDWRWLQLAVSLPTFLFLLYYWFVPES 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         50 --WVHMLDNN-----TGSGNETGILSEDALLRISIPLDSNLRPEKCRRFVHPQWQLLHLN 102
                                                                                                                                                                                                                                                                                                                        175 ARWLIITNKLDEGLKALRKVARTNG-IKNAEETLNIEVVRSTMQEELDAAQTKT-TVCDL 232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 DDVLBQVGBFGWFQXQAFLLLCLISASLAPIYVGIVFLGFT---PGHYCQNPGVAELSQR 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20005-3315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 12.1%; Score 169.5; DB 3; Length 556; Similarity 20.5%; Pred. No. 8.9e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels 157; Gaps
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STREET:
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US-08-501-572-2
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APPLICANT: Grundeman, Dirk
APPLICANT: Grundeman, Dirk
APPLICANT: Grundeman, Dirk
TITLE OF INVENTION: Transport protein Which Effects The
TITLE OF INVENTION: Transport Of Cationic Xenobiotics and\or Pharmaceuticals,
NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 10.8%; Score 152.5; DB 3; Length 555; Best Local Similarity 19.8%; Pred. No. 8e-09; Matches 71; Conservative 40; Mismatches 109; Indels 139; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             67 LSEDALLRISIP---LDSNLRPEKÇRRFVHPOMQLLHLNGTIHSTSEADTE-----PÇ 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65 WSPAEELNYTVPGPGPAGEASPRQCRRY-EVDWNQSTFD-CVDPLASLDTNRSRLPLGPC 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            117 VPGWVYDQSYFPSTIVTKWDLYÇDYQSLKSVVQFLLLTGMLVGGIIGGHVSDR------ 169
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| SSIVTEFNLVCANSWMLDLFQSSVNVGFFIGSMSIGYIADRFGRKLCL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 GGLGRFQMLHLVFILPSLMLLIPHILLE-NFAARIPGHRC---WHMLDNNTGSGNETGI 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12 GGEFHFFQKQMFFLLALLSATFAPİYVGIVFLGFTÞDHRCRSPGVAELSIRCG----- 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -----ESARWLITNKLDEGL 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 YQVAYTVGLLVLAGVAYALPHWRWLQFTVALPNFFFLLYYWCIPESPRWLISQNKNAEAM 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            189 KALRKVARTNGIKNAEETLINIEVVRSTMOEELDAAQTKTTVCDLFRNPSMFKRICTLVF 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & Dunner STREET: 1300 I Street, N.W., Suite 700
CITY: Washington
                                                                                           ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIOW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 02481.1453-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)408-4000
TELEFAX: (202)408-4400
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                               APPLICATION NUMBER: US/08/501,572
                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Toohey, Kimberlin M
REGISTRATION NUMBER: 35,391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/09040444
Patent No. 6063766
GENERAL INFORMATION;
                                                                                                                                                                                                                                                                                                                                                                                                                     s 555 amino acids amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: peptide US-08-501-572-3
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                                                                                                                                                                                                                                                  CLASSIFICATION:
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                                                                                                                                                                                                                                    FILING DATE:
                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
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Sequence 2, Application US/08501572
| Patent No. 6053623
| GENERAL INFORMATION. | APPLICANT: Koepsell, Hermann | APPLICANT: Grundeman, Dirk | APPLICANT: Groboulev, Valentin | TITLE OF INVENTION: Transportr protein Which Effects The | TITLE OF INVENTION: Transportr protein Which Effects The | TITLE OF INVENTION: Transport Of Cationic Xenobiotics and/or Pharmaceuticals, ITTLE OF INVENTION: DNA Sequences Encoding It And Their Use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65 WSPARELNYTVPGPGPAGEASPROCRRY-EVDWNOSTFD-CVDPLASLDTNRSRLPLGPC 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ||||||:
| SSIVTEFNLVCANSWMLDLFQSSVNVGFFIGSMSIGYIADRFGRKLCL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               170 ----- 172
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                       ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P.
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                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: O'CONDOT, Steven P
REGISTRATION NUMBER: 41,225
REFERENCE/DOCKET NUMBER: 2481.1453-01
TELEPHONE: (202) 408-4400
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                STATE:
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
"""FR: IBM PC compatible
"""FR: IRM PC compatible
                                                                                                                                                                                                                                                                                             US/09/040,444
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/04
FILING DATE: March 18, 1998
CLASSIFICATION:
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TYPE: amino acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS:
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MOLECULE TYPE: peptide
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US-08-501-572-2
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                   US-09-040-444-2
                                                                                                                   RESULT 9
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER: 35,391
REFERENCE/DOCKET NUMBER: 02481.1453-00000
TELECOMMUNICATION INFORMATION:
GENERAL INFORMATION:
APPLICANT: Koepsel.
APPLICANT: Grundem.
APPLICANT: Gorboul
                                                               Sequence 2, Application US/09040444 patent No. 6063766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1300 I St
CITY: Washington
STATE: D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/501,572
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                                                                                                                                                                                                                                                                                                                                                                                                                                               114 SHLPLGÞÖQDGWVYDTP--GSSÍVTEFNLVÖADSWKLDLFQSCLNAGFFFGSLGVGYFAÐ 171
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                                                                                                                                                                                                                                                                                                           232 SRRTVAIMYOMAFTVGLVALTGLAYALPHWRWLQLAVSLPTFLFLLYYWCVPĖSPRWLLS 291
                                                                                                                                                                                                                                                                                                                                                170 -----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 GNETGILSEDALLRISIPLDSNLRPE-----KCRREVHPQWQLLHLN------GTIHST 108
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                                                                                                                                                                                                                                         292 OKRNTEAIKIMDHIAQKNG-KLPPADLKMLSLEEDVTEKL----SPSFADLFRTPRLRK 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    64 -----WSPAEELNYTVP---GLGPAGEAFLGQCRRY-EVDWNQSALSCVDPLASLATUR 113
                                                                                                                                                                                                         241 RICILVFL 248
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                                                                                                                                                                        346 RTFILMYL 353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEADIEPCVPGWYYPOSYFPSTIVIKWDLYCDYQSLKSVVQFLLLTGMLVGGIIGGHVSD 168
                                                                                                                                                                                                                                                                                                                                                                                  REGRKLCLLGTVLVNAVSGVLMAESPNYMSMLLERLLQGLVSKGNMMAGYTLITEEVGSG 231
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch 10.4%; Score 146; DB 3; Length 553;
1 Similarity 20.4%; Pred. No. 4.5e-08;
75; Conservative 34; Mismatches 117; Indels 142; Gaps
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1300 I Street, N.W., Suite 700
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   Koepsell, Hermann
Grundeman, Dirk
Gorboulev, Valentin
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US-09-040-444-2
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US-08-964-127-2
; Sequence 2, Application US/08964127
; Patent No. 6277565
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                                                       RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Transport protein which Effects The TITLE OF INVENTION: Transport Of Cationic Xenobiotics and or Pharmaceuticals, TITLE OF INVENTION: DNA Sequences Encoding It And Their Use.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 41,225
REFERENCE/DOCKET NUMBER: 245
TELECOMMUNICATION INFORMATION:
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STREET: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC comparible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
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                                                                                                                                                                             292 OKRNTEAIKIMDHIAOKNG-KLPPADLKMLSLEEDVTEKL----SPSFADLFRTFRLRK 345
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                                                                                                                                             241 RICILVFL 248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ch 10.4%; Score 146; DB 3; Length 553; 1 Similarity 20.4%; Pred. No. 4.5e-08; 1 Similarity 20.4%; Mismatches 117; Indels 142; 75; Conservative 34; Mismatches 117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
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99 LHLNGTIHSTSEADTEPÇVDGWVYD-QSYFPSTIVTKWDLYCD--YQSLKSVVQFLL--- 152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 HYGAFPPNASGW-EQPPNASGVSVASAALAASAASKVATSTDPS-----CSGFAPP---- 65
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1 Similarity 18.2%; Pred. No. 0.0028;
64; Conservative 33; Mismatches 92; Indels 162;
                 APPLICANT: Grandearl, Andrew David John
TITLE OF INVENTION: NOVEL GENES ENCODING TRANSPORTER-LIKE
TITLE OF INVENTION: MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         283 NTCPLPATSSFSFASLLNYRN--IWRNLLILGFINFIAHAIRH---CYOPV 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  228 TVCDL-----FRNPSMRKRICILVFLRKKISRKRHKNDCYTKY 265
                                                                                                                                                                             ZOUNTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: 1 DBA Compatible
COPERATING SYSTEM: Windows 95
SOFTWARE: FastEND for Windows Version 2.0b
APPLICATION NUMBER: US/08/964,127
FRING APPLICATION DATA:
FRING APPLICATION DATA:
FRING APPLICATION DATA:
FRING APPLICATION DATA:
FRING APPLICATION NUMBER: US/08/964,127
FRIOM TOWNER: US/08/964,127
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ATTORNEY/AGENT INFORMATION:
NAME: Crews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
REFERENCE/DOCKET NUMBER: 07334/038001
TELECOMMUNICATION INFORMATION:
                                                                                                    E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO. 2: SEQUENCE CHARACTERISTICS:
LENGTH: 520 ami-
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617/542-8906
                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
FRAGMENT TYPE: internal
                                                                 NUMBER OF SEQUENCES:
GENERAL INFORMATION:
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                                                                                                                                      Boston
                                                                                                                                                                       USA
                                                                                                                                                                    COUNTRY:
                                                                                                                  STREET:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 HYGAFPPNASGW-EQPPNASGVSVASAALAASAASRVATSTDPS-----CSGFAPP---- 65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
7.4%; Score 104; DB 4; Length 520;
Best Local Similarity 18.2%; Pred. No. 0.0028;
Matches 64; Conservative 33; Mismatches 92; Indels 162;
                     PEPLICANT: Grandearl, Andrew David John
TITLE OF INVENTION: NOVEL GENES ENCODING TRANSPORTER-LIKE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
GITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  228 TVCDL-----FRAPSMRKRICILVFLRKKISRKRHKNDCYTKY 265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        162 -----VSDRW------
                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastESQ for Windows Version 2.0b
APPLICATION NUMBER: US/09/496,692
                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Crews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
REFERENCE/DOCKET NUMBER: 07334/038001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/964,127
FILING DATE: 06-NOV-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 617/542-8000
TELEX: 200154
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 520 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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FRAGMENT TYPE: internal
US-09-496-692-2
GENERAL INFORMATION:
                                                                                                                                                                                                       02110-2804
                                                                                                                                                                                         USA
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                                                                                                                                                                                       COUNTRY:
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US-08-928-692-12
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; Sequence 12, Application US/08928692; Patent No. 5958727

; Sequence 2, Application US/09496692; Patent No. 6313271

US-09-496-692-2

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US-08-928-692-12
                                                                                                                                                                                                                                                                                          US-09-339-972-12
                                                                                                                                                                                                                                                                                                                  RESULT 13
                                                                                                                                                                                                                      Sequence 12, Application US/09339972
Patent No. 6323002
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 7.1%; Score 100; DB 2; Length 524; Best Local Similarity 27.6%; Pred. No. 0.0082; Matches 35; Conservative 22; Mismatches 34; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 212-878-9655
NFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
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LENGTH: 524 amino acid
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FILING DATE: 12-SEPT-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Elias J
REGISTRATION NUMBER: 33,728
REFERENCE/DOCKET NUMBER: 4944.200-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
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                                                        TITLE OF INVENTION: Methods for MCTITLE OF INVENTION: a POLYPEPTIDE NUMBER OF SEQUENCES: 80 CORRESPONDENCE ADDRESS:
                                                                                                                                       APPLICANT: Brody, Howard
APPLICANT: Yaver, Deborah S.
APPLICANT: Lames, Michael
APPLICANT: Hansen, Kim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                               304 ILVALML 310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        184 LDEGLKALRKVARTNGIKNABETINIEVVRSTMQEELDAA--QTKTTVCDLFRNPSMRKR 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 193 QLAIVTGILISQIIGLEFILGNY-DLWHILLGLSGVRAILQSLLLFFCPESPRYLYI--K 249
                                                                                                                                                                                                                                                                                                                                                                                                                    242 ICILVFL 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          149 QFLLLTGMLVGGIIG-----GHVSDRWLV---------ESARWLIITNK 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: sir
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                   STREET:
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New York
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                 405 Lexington Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hansen, Kim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brody,
Yaver,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                       No. 6323002o No. 6323002disk of No. 6323002th America,
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                                                                                                                      Methods for Modifying the Production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80
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                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/08964127
Patent No. 6277565
GENERAL INFORMATION:
APPLICANT: Grandearl, Andrew Dav
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 7.1%; Score 100; DB 4; Length 524; Best Local Similarity 27.6%; Pred. No. 0.0082; Matches 35; Conservative 22; Mismatches 34; Indels
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                 COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/964,127
FILING DATE: 06-NOV-1997
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TELECOMMUNICATION INFORMATION:
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NAME: Lambiris, Elias J
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: No. 6323002e
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                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   184 LDEGLKALRKVARTNGIKNAEETLNIEVVRSTMQEELDAA--QTKTTVCDLFRNPSMRKR 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              193 QLAIVTĞILISQIIGLEFILGNY-DLWHILLGLSGVRAILQSLLLFFCPESPRYLYI--K 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      149 OFLLLTGMLVGGIIG-----GHVSDRWLV-------ESARWLIITNK 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              242 ICILVEL 248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                              ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
                                                                                                                                                                                            02110-2804
                                                                                                                                                                                                                                                          Boston
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                                                                                                                                                                                                                  USA
                                                                                                                                                                                                                                                                                                                                                                                        Grandearl, Andrew David John
VENTION: NOVEL GENES ENCODING TRANSPORTER-LIKE
                                                                                                  IBM Compatible
SYSTEM: Windows 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM Compatible
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NER: 4944.200-US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     171 ---LVESARWLIITNKLDEGLKALRKVA---RTNGIKNAEETLNIEVVRSTMQEELDAAQ 224
                                                                                                                                                                                                                                                                                                                                                                                                Query Match 7.0%; Score 98.5; DB 4; Length 286;
Best Local Similarity 17.8%; Pred. No. 0.0049;
Matches 54; Conservative 29; Mismatches 73; Indels 147; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 HYGAFPPNASGW-EQPPNASGVSVASAALAASAASRVATSTDPS-----CSGFAPP---- 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 39 NFAAAIPGHRCWVHMLDNNTGSGNETGILSEDALLRISIPLDSNLRPEKCRRFVHPQWQL 98
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Sequence 4, Application US/09496692

Sequence 4, Application US/09496692

Patent No. 6313271

GENERAL INFORMATION:

APPLICANT: Grandearl, Andrew David John on TITLE OF INVENTION:

MUMBER OF INVENTION:

NUMBER OF SEQUENCES: 17

CORRESPONDENCES: 17

STREET: 225 Franklin Street

CITY: Boston
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMpatible
OPREATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/496,692
                                                        NAME: Crews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
REFERENCE/DOCKET NUMBER: 07334/038001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
                                                                                                                                                            TELEFAX: 617/5/7-2-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS: LENGTH: 286 amino acide TYPE: amino acide TOPOLOGY: linear
                   FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein FRAGMENT TYPE: internal
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99 LHLNGTIHSTSEADTEPCVDGWVYD-QSYFPSTIVTKWDLVCD--YQSLKSVVQFLL--- 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -----LTGMLVGGI---- 161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ----OEALODLE 282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 7.0%; Score 98.5; DB 4; Length 286; Best Local Similarity 17.8%; Pred. No. 0.0049; Matches 54; Conservative 29; Mismatches 73; Indels 147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Crews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
REFERENCE/DOCKET NUMBER: 07334/038001
TELECOMMUICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: January 6, 2003, 22:51:39
Job time : 23 secs
FILING DATE: 06-NOV-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 200154
INFORMATION FOR SEQ 1D NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 286 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX:
TELEX: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-496-692-4
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OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         January 6, 2003, 22:48:58; Search time 16 Seconds (without alignments) 317.438 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MAFBELLSQVGGLGRFQMLH......RKKISRKRHKNDCYTKVTKF 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    117078 segs, 18951520 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Published_Applications_AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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| 90.5 90 89.5 83.5 | 90.55 90.55 | 551 344 216.5 193.5 174 160 | Score 600 592 |
| 0000 444 44410 | 7.000 00.00 00444 | 39.2 24.5 15.4 12.4 11.4 | Query Match Length |
| 535 547 488 494 | 286 81 486 487 | 553 113 557 557 101 44 | ength I |
| 10 12 10 | 10 10 10 | 9 9 10 10 10 10 10 10 10 10 10 10 10 10 10 | 9 9 8d |
| US-09-795-693-20 US-10-155-891-2 US-10-094-059-4 US-09-349-015-35 US-09-381-947A-5 | US-10-000-273-4 US-09-864-761-46676 US-09-860-232A-7 US-10-095-139-14 US-09-795-693-27 | US-10-03-13-17 US-09-864-761-42488 US-09-798-743A-3 US-09-798-743A-1 US-10-050-786-8 US-10-095-139-15 US-10-005-273-2 | ID |
| Sequence 2, Appli Sequence 4, Appli Sequence 35, Appli Sequence 35, Appli Sequence 5, Appli | Sequence 4, Appli Sequence 46676, A Sequence 7, Appli Sequence 14, Appl Sequence 27, Appl | 2, 13 6 13 7 14 7 15 7 | Description Sequence 5, Appli Sequence 16, Appli Sequence 17, Appl |

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|------------------|-------------------|-------------------|------------------|------------------|-----------------|----------------|---------------------|-------------------|-----------------|-------------------|-----------------|--------------------|--------------------|--------------------|---------------------|------------------|-----------------|------------------|-----------------|---|---------------------|-------------------|-------------------|-------------------|-------------------|-------------|-------------|--------------------|-----------------|-------------------|-----|
| 75 | 10.0 | 10.0 | 75.5 | 76 | 76 | 76 | 6.0 | 1 | 76.5 | 78 | 78.5 | 79 | 79 | 80.5 | | о () | 80.5 | 80.5 | 80.5 | 81.5 | α - | - 0 | 8 | 82 | 82 | 82 | 22 | | מ | 83 | |
| υ. | | л (| л | 5.4 | 5.4 | 5.4 | | | л | თ თ | 5. 6 | 5.6 | 5.6 | | 1 (| 7 | 5.7 | 5.7 | 5.7 | υ· α | | | | | 5.8 | | | | | 5.9 | |
| 526 | | 436 | 209 | 1203 | 1203 | 314 | , i | 1648 | 1642 | 574 | 476 | 664 | 000 | 1 4 | 100 | 325 | 325 | 325 | 323 | 5 6 5 | 1 (| л Э | 501 | 471 | 471 | Ų Ų | . d | | 572 | 406 | |
| v | , | 10 | ø | φ | v | 5 | , , | - | 10 | 10 | v | , [| ; = | . (| 0 | 10 | 10 | 10 | 5 | , (| ٠, | ٥ | 10 | 10 | 10 | - | , , | 2 | 10 | 10 | |
| 02-10-024-020-20 | 10 004 603 30 | US-09-922-501-8 | US-10-108-915-30 | US-09-909-280A-2 | OS-09-990-040-4 | CG-CG-CG CAE-3 | 110 00 067-550-1040 | TIS-09-925-442-35 | US-09-925-442-2 | US-09-225-502-6 | 08+10+155-631-4 | OS-09-813-242-1217 | US-09-015-242-0179 | TO 00-015-040-5470 | 110-09-738-626-4277 | US-09-804-006-10 | US-09-739-451-5 | US-09-804-35/-IO | US-09-9/4-914-0 | CO + C - CO + C - C - C - C - C - C - C - C - C - C | 110 10 101 1010 104 | 115-10-051-902-24 | US-09-778-927A-48 | US-09-778-92/A-49 | US-09-7/8-92/A-47 | -09-770-027 | 00 779-9774 | 115-09-778-927A-51 | US-09-919-781-2 | US-09-925-29/-/26 | , |
| | Seguence 30, App⊥ | Sequence 8, Appli | 0 | | v. | N | Seguence 1240, Ap | Sequence 35, Appr | | comicance 2 Appli | ָ ס | | | - | Sequence 4277, AP | | 3 , | | 5. | | Sequence 24, Appl | Sequence 24, Appl | , , | ָ מ | | 47 | 50 | Sequence 51, Appl | :: | | 726 |

ALIGNMENTS

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RESULT 1
US-10-095-139-5
US-10-095-139-5
Sequence 5, Application US/10095139
Patent No. US20020163357A1
JETUE NO. US200201631A
JETUE NO. US2010-03-11
JETUE NO. US2010-03-12
PRIOR FILING DATE: 2002-03-11
JETUE APPLICATION NUMBER: US/10/095,139
CURRENT FILING DATE: 2002-03-11
JETUE APPLICATION NUMBER: US/10/095,139
CURRENT FILING DATE: 2002-03-12
PRIOR FILING DATE: 2002-03-12
PRIOR FILING DATE: 2002-03-12
JETUE PRI
JETUE APPLICATION NUMBER: US/10/095,139
CURRENT APPLICATION NUMBER: US/10/095,139
CURRENT APPLICATION NUMBER: US/10/095,139
JETUE PRI
JETUE NOS: 24
NUMBER OF SEQ ID NOS: 24
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APPLICANT: Silos-Santiago, Inmaculada APPLICANT: Millennium Pharmaceuticals, Inc. TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic TITLE OF INVENTION: 1on Transporters and Uses Therefor FILE REFERENCE: MPIO1-017PIRNM

CURRENT APPLICATION NUMBER: US/10/095,139
CURRENT FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: 60/275,172
PRIOR FILING DATE: 2001-03-12
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FREESEQ for Windows Version 4.0

SEQ ID NO 17

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                                                          181 LOMAVMGTAAAFAPAFPVYXXXRFLLAFHASGAEPLGLLAVMEWTAARARPLVMTLNSLG 240
                                                                                          ------WLVESARWLITNKLDEGLKALR 192
                                                                                                           241 FSFGHGLTAAVAYGVRDWTLLQLVVSVPFFLCFLYSWWLAESARWLLTTGRLDWGLQELW 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          59 GSGNETGILSEDALLRISIPLDSNLRPEKÇRRFVHPQWQLLHLNGTIHSTSEADTEPCVD 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  57 GSAVSTN-MTPKALLTISIPPGPNQGPHQCRRPRQPQWQLLDPNATATSWSEADTEPCVD 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   119 GWVYDQSYFBSTIVTKWDLVCDYQSLKSVVQFLLLTGMLVGGIJGGHVSDR----- 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
42.1%; Score 592; DB 9; Length 550;
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Matches 143; Conservative 28; Mismatches 69; Indels 116; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          176 CCLQLAVAGTSTIFAPTFVIYCGLRFVAAFGMAGIFLSSLTLMVEWTTTSRRAVTMTVVG 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ------WLVESARWLIITNKLDEGLKA 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   236 CAFSAGQAALGGLAFALRDWRTLQLAASVPFFAISLISWWLPESARWLIIKGKPDQALQE 295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MAFEELLSQVGGLGRFQMLH-LVFILPSLMLLIP-HILLENFAAAIPGHRCWVHMLDNNT 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 191 LRKVARTNGIKNAEETLNIEVVRSTMOEELDAAQTKTTVCDLFRNPSMRKRICILV 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   193 KVARTNGIKNAEETINIEVVRSTMQEELDAAGTKTTVCDLFRNPSMRKRICI 244
                                                                                                                                                                   301 RVAPINGKGAVQDTLTPBVLLSAMREELSMGQPPASLGTLLRMPGLRFRTCI 352
                                                                                                                                                                                                                                                                              Sequence 16, Application US/10095139
Parent No. US20020165357A1
                                                                                          170 -----
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ORGANISM: homo sapiens
                               170 -----
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US-10-095-139-16
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Sequence 17, Application US/10095139; Patent No. US20020165357A1; GENERAL INFORMATION: APPLICANT: Curtis, Rory A.J.

US-10-095-139-17

RESULT 3

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**PAPELICANT: Penn, Sharron G.
**APPLICANT: Penn, Barron G.
**APPLICANT: Penn, David R.
**APPLICANT: Hanzel, David R.
**APPLICANT: Hanzel, David K.
**APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: HUMAN GENE EXPRESSION ANALYSIS BY MICROARRAY
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
TITLE NEPERBNUE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR PILING DATE: 2000-05-26
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                                                                                                          n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  181 LLVSVSGTAAAFMFTFPLYCLFRFLLASAVAGVMMNTASLLMEWTSAQGSPLVWTLNALG 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       170 -----WLVESARWLIITNKLDEGLKALR 192
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                                                                                                                                                                                                                       1 MAFBELLSQVGGLGRPQMLHLVFILPSLMLLIPHILLENFAAAIPGHRCWVHMLDNNTGS 60
                                                                                                                                                                                                                                                           193 KVARTNGIKNAEETLNIEVVRSTMOEELDAAQTKTTVCDLFRNPSMRKRICI 244
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                                                                                                                                                               Matches 120; Conservative
TYPE: PRT
ORGANISM: Mus musculus
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                                                        US-10-095-139-17
                                                                                                             Query Match
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US-09-864-761-42488
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 42488
SEQ ID NO 42488
                                                                                                                                                                                                                                                                                                                                                                                          US-09-798-743A-3
                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/09798743A Patent No. US20020099093A1 GENERAL INFORMATION:
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OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 7.2
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 5.7
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 7
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FEART, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: SWISSPROT HIT: Q920E8, EVALUE 3.00e-05
OTHER INFORMATION: EST_HUMAN HIT: AV652088.1, EVALUE 4.00e-32
                        APPLICANT: Nezu, Jun-ichi
APPLICANT: Ose, Asska
APPLICANT: Ose, Asska
APPLICANT: Ose, Asska
CORRENT: Ose, ASSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
FILE REFERENCE: 06501-073001
CURRENT APPLICATION NUMBER: US/09/798,743A
CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: PCT/JP99/04853
PRIOR FILING DATE: 1999-09-07
PRIOR FILING DATE: 1999-09-07
NUMBER OF SEQ ID NOS: 31
NUMBER OF SEQ ID NOS: 31
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OTHER IN
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SOFTWARE: PatentIn Ver. 2.0
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63 RPHQCRRPROPOWOLLDPNATATSWSEADTEPCVDGWVYDRSIFTSTIVAK 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  84 RPEKCRRFVHPQWQLLHLNGTIHSTSEADTEPCVDGWVYDQSYFPSTIVTK 134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 24.5%; Score 344; DB 10; Length 113; Local Similarity 59.5%; Pred. No. 3.1e-28; es 66; Conservative 12; Mismatches 33; Indels
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00661
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LENGTH: 557
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APPLICANT: Nezu, Jun-ichi
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                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: PCT/JP99/04853
PRIOR FILING DATE: 1999-09-07
PRIOR APPLICATION NUMBER: JP 10-252683
PRIOR FILING DATE: 1998-09-07
PRIOR FILING DATE: 1998-09-07
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CURRENT FILING DATE: 2001-03-02
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TITLE OF INVENTION: SYSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
FILE REFERENCE: 06501-073001
                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 31
                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                  LENGTH: 557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             157 SFISGQLSDRFGRKNVLFLTMGMQTGFSFLQVFSVNFEMFTVLFVLVGMGQISNYVAAFV 216
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Local Similarity 23.4%; Pred. No. 3.3e-14;
Les 91; Conservative 34; Mismatches 89; Indels 175.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       217 LGTEILSKSIRIIFATLGVCIFYAFGFMVLPLFAYFIRDWRMLLLALTVPGVLCGALWWF 276
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                                                                                                                                                                      / Match 13.8%; Score 193.5; DB 10; Length 557;
Local Similarity 22.0%; Pred. No. 7.6e-12;
Nes 83; Conservative 37; Mismatches 106; Indels 151; Gaps
                                     59 GSGNETGILSEDALLRISIPL---DSNLRPEKCRRF---VHPQWQLLHLN-GTIHSTSEA 111
61 AWRNHT---
                                                                                                                             3 FEELLSQVGGLGRPQMLHLVFILPSLMLLIPH----ILLENFAAAIPGHRCWVHMLDN-NT 58
                                                                                   4 YDEVTAFLGEWGPFQ--RLIFFILSAS-IIPNGFTGLSSVFLIATPEHRCRVPDAANLSS 60
  ----VPLRLRDGREVPHSCRRYRLATIANFSALGLEPGRDVDLGQL 108
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92; Indels 162; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Grandearl, Andrew David John
TITLE OF INVENTION: NOVEL GENES ENCODING TRANSPORTER-LIKE
MOLECULES
                                                                                                                                                                                                                                                                  Length 44;
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7.4%; Score 104; DB 9; Length 520;
Best Local Similarity 18.2%; Pred. No. 0.011;
Matches 64; Conservative 33; Mismatches 92; Indels 1
                                                                                                                                                                                                                                                                                                        8; Indels
                                                                                                                                                                                                                                                                                                                                         102 NGTIHSTSBADTEPCVDGWVYDQSYFPSTIVTKWDLVCDYQSLK 145
                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IDM Compartible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
RILING DATE: 02-No. US20020160386A1-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                        Query Match
11.4%; Score 160; DB 9;
Best Local Similarity 63.6%; Pred. No. 6.4e-10;
Matches 28; Conservative 8; Mismatches 8;
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FILING DATE: 06-NOV-1997
ATTORNEY/AGENT INFORMATION:
NAME: Crews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-6070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                     NUMBER OF SEQ ID NOS: 24
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15
LENGTH: 44
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MOLECULE TYPE: protein
FRAGMENT TYPE: internal
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/10000273; Patent No. US20020160386Al GENERAL INFORMATION:
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                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
    PRIOR FILING DATE: 2001-03-12
                                                                                                                                                                                 ; OTHER INFORMATION: consensus
US-10-095-139-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 200154
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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ZIP: 02110-2804
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                                                                                                                                                              FEATURE:
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Patent No. US20020155339A1

GENERAL INFORMATION:

APPLICANT: BUDDE et al.

TITLE OF INVERNATION:

FILE REPERENCE: PT0137P1C1

CURRENT APPLICATION NUMBER: US/10/050,786

CURRENT APPLICATION NUMBER: US/10/050,786

CURRENT PILING DATE: 2002-01-18

PRIOR APPLICATION NUMBER: US 09/774,028

PRIOR APPLICATION NUMBER: US 60/145,958

PRIOR FILING DATE: 1000-07-27

PRIOR FILING DATE: 1999-00-18

PRIOR FILING DATE: 1999-00-18

PRIOR FILING DATE: 1999-00-18

PRIOR FILING DATE: 1999-00-18

PRIOR FILING DATE: 1999-01-18

PRIOR FILING DATE: 1999-03-18

PRIOR FILING DATE: 1999-03-14

NUMBER OF SEQ ID NOS: 12

SOFTWARE: PATENTIN VET. 3.1
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112 DTEPCVDGWVXDQSYFPSTIVTKWDLVCDYQSLKSVVQFLLLTGMLVGGIIGGHVSDR-- 169
                              169 RKDVJ.FVTMGMQTGFSFLQIFSKNFEMFVVLFVLVGMGQISNYVAAFVLGTEILGKSVRI 228
                                                                                                                                                                                           .-----WLV-ESARWLIITN 182
                                                                                                                                                                                                                            229 IFSTLGVCIFYAFGYMVLPLFAYFIRDWRMLLVALTMPGVLCVALWWFIPESPRWLISQG 288
                                                                                                                                                                                                                                                                    183 KLDEGLKALRKVARTNGIKNAEETLNIEVVRST-----MQEELDAAQTKTTVCDLFRNP 236
                                                                                                                                                                                                                                                                                                               Gaps
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12.4%; Score 174; DB 9; Length 101;
Best Local Similarity 100.0%; Pred. No. 7.3e-11;
Matches 34; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Curtis, Rory A.J.
APPLICANT: Curtis, Rory A.J.
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Millennium Pharmaceuticals, Inc.
TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic
TITLE OF INVENTION: Ion Transporters and Uses Therefor
FILE REFERENCE: MPIOJ. 017PARNM
CURRENT APPLICATION NUMBER: US/10/095,139
CURRENT FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: 60/275,172
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Patent No. US20020165357A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                    289 RFEEAEVIIRKAAKANGI---
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ORGANISM: Homo sapiens
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SEQ ID NO 8
LENGTH: 101
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US-10-095-139-15
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| FILING DATE: 2001-01-29 R OF SEQ ID NOS: 49117 | QY 39 NFAAAIPGHKCWVHMLINNIGSGNEIGILEEDHALERISIEDDSHIREENCESCH 15 1 1 1 1 1 1 1 1 1 |
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| FRIOR APPLICATION NUMBER: US PRIOR APPLICATION UNMBER: US PRIOR APPLICATION UNMBER: US | atch 7.0%; Score 98.5; DB 9; Length 286; cal Similarity 17.8%; Pred. No. 0.017; cal Conservative 29; Mismatches 73; Indels 147; Ga |
| APPLICATION N FILING DATE: APPLICATION N FILING DATE: | 2Q ID NO: 4: |
| PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT/US01/00662 PRIOR FILING DATE: 2001-01-30 | TH: 2 : ami LOGY: |
| PRIOR FILING DATE: 2001-01-30 PRIOR FILING DATE: 2001-01-30 PRIOR FILING DATE: 2001-01-30 | ; TELEX: 200154 ; INFORMATION FOR SEQ ID NO: 4: ; SEQUENCE CHARACTERISTICS: |
| ; PRIOR APPLICATION NUMBER: PCI/USO1/00009; PRIOR FILING DATE: 2001-01-30 ; PRIOR APPLICATION NUMBER: PCT/USO1/00665 ; PRIOR ETIING DATE: 2001-01-30 | TION 5070 |
| ; PRIOR FILING DATE: 2001-01-30 ; PRIOR APPLICATION NUMBER: PCT/US01/00664 ; PRIOR FILING DATE: 2001-01-30 ; PRIOR FRIITGATION NUMBER: PCT/US01/00669 | NAME: Crews, Ph.D., L. Lee REGISTRATION NUMBER: P-43,567 REFERENCE/DOCKET NUMBER: 07334/038001 |
| PRIOR APPLICATION NUMBER: PCT/US01/00666 PRIOR FILING DATE: 2001-01-03 PRIOR APPLICATION NUMBER: PCT/US01/00667 | FILING DATE: US/08/964,127 FILING DATE: 06-NOV-1997 ATTORNEY/AGENT INFORMATION: |
| PRIOR FILING DATE: 2000-10-04 PRIOR FILING DATE: 2000-09-27 PRIOR FILING DATE: 2000-09-27 | APPLICATION NUMBER: US/10/000,273 FILING DATE: 02-No. US20020160386A1-2001 PRICE NUMBER: USATA: |
| FILING DATE: 2000-08-03 APPLICATION NUMBER: GB | , OPERATING SYSTEM: Windows 95 ; SOFTWARE: FastSeQ for Windows Version 2.0b ; CURRENT APPLICATION DATA: |
| | IBM Compatible |
| | VIRY: USA: 02110-28 |
| REFERENCE: Acomi | ; STREET: 225 Franklin Street ; CITY: Boston STATE: MA |
| ANT: Chen, OF INVENTIO | |
| שני שני ש | APPLICANT: GRANGEST, ANGLES ENCODING TRANSPORTER-LIKE TITLE OF INVENTION: NOVEL GENES ENCODING TRANSPORTER-LIKE MOLECULES |
| RESULT 11 US-09-864-761-46676 US-09-864-761-46676, Application US/09864761 ; Sequence 46676, Application US/09864761 ; Patent No. US20020048763A1 ; GENERAL INFORMATION: | RESULT 10 US-10-000-273-4 ; Sequence 4, Application US/10000273 ; Patent No. US20020160386A1 ; Patent INFORMATION: |
| Db 283 SST 285 | 883 NICFHEAISSESEASHINIRNIMANHHIHDE INE FRISTANI |
| Qy 225 TKT 227 | 228 |
| Qy 171LVESARWLIITNKLDEGLKALRKVARTNGIKNA | QY 171 |
| Db 173 YLMRLELCDPTQRLRVALAGELVGVGGHFLFLGLALVSK | 173 Y |
| 162 | Qy 162 170 |
| Db 113 SGYLFLGYPADRFGRRGIVLLTLGLVGPCGVGGAAAGSS | |
| יי ווי ווי ווי ווי ווי ווי ווי ווי ווי | 153 |
| QY 99 LHLNGTIHSTSEADTEPCVDGWVYD-QSYFPSTIVTKWD | Qy 99 LHLNGTIHSTSEADTEPCVDGWVYD-QSYFPSTIVTKWDLVCDYQSLKSVVQFLL 152 b 66bfnhclkdwdynglpvlttnaigqwdlvcdlgwqvileqilfilgfa 112 |

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                                                                                                                                                CTHER INFORMATION: MAP TO AB026898.1
CTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN LUNG. SIGNAL = 1
OTHER INFORMATION: EST HUMAN HIT: AV652088.1, EVALUE 3.10e-02
CTHER INFORMATION: SWISSPROT HIT: Q62674, EVALUE 6.20e+00
US-09-864-761-46676
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Patent No. US20020165357A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Milennium Pharmaceuticals, Inc.
TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic
TITLE OF INVENTION: Lon Transporters and Uses Therefor
FILE REFERENCE: MPI01-017PIRNM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/09860232A
Patent No. US20020028494A1
GENERAL INFORMATION:
APPLICANT: CURTIS, ROLY A.J.
TITLE OF INVENTION: 57256 AND 58289, NOVEL HUMAN
TITLE OF INVENTION: TRANSPORTERS AND USES THEREOF
FILE REPERENCE: 381552001500
CURRENT APPLICATION NUMBER: US/09/860,232A
CURRENT FILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/205,288
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Consensus amino acid sequence US-09-860-232A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             107 STSEADTEPCVDGWVYDQSYFPS 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                     SEQ ID NO 46676
LENGTH: 81
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US-09-860-232A-7
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US-10-095-139-14
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144 LKSVVQFLLLTGMLVGGIIGGHVSDRWLVESARWLIITNKLDEGLKALRKVARTNGIKNA 203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Glucksmann, Maria A. TILL4, 23821, 33894, and TITLE OF INVENTION: 32613, No. US20020068710Alel Human Transporters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Glucksmann, Maria A.
TITLE OF INVENTION: 20685, 579, 17114, 23821, 33884, and
TITLE OF INVENTION: 32613, No. US20020068710A1e1 Human Transporters
FILE REPERENCE: 35800/209292
CURRENT APPLICATION NUMBER: US/09/795,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
6.4%; Score 90.5; DB 9; Length 487;
8est Local Similarity 26.4%; Pred. No. 0.23;
Matches 29; Conservative 28; Mismatches 32; Indels 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    204 BETLNIEVVRSTMQEELDAAQT-KTTVCDLFR---NPSMRKRICILVFLR 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            33 DQ--EIQEIKAELEAGVEEEKAGKASWGELFRGRTRPKVRQRLLMGVMLQ 282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Pfam consensus sequence US-09-795-693-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 38800/209292
CURRENT APPLICATION NUMBER: US/09/795,693
CURRENT FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/185,906
PRIOR PILING DATE: 2000-02-29
NUMBER OF SEQ ID NOS: 42
SOFTWARE: FastSEQ for Windows Version 4.0
CURRENT APPLICATION NUMBER: US/10/095,139
CURRENT FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: 60/275,172
PRIOR FILING DATE: 2001-03-12
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASEUSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 27, Application US/09795693; Patent No. US20020068710A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: consensus
US-10-095-139-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-795-693-27
                                                                                                                                                                      SEQ ID NO 14
LENGTH: 487
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                                                                                                                                                                                                                                                                                                    FEATURE:
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CURRENT FILING DATE: 2001-02-28

PRIOR APPLICATION NUMBER: 60/185,906

PRIOR APPLICATION NUMBER: 60/185,906

PRIOR FILING DATE: 2000-02-29

NUMBER OF SEQ ID NOS: 42

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 20

LENGTH: 535

TYPE: PRT

ORGANISM: Artificial Sequence

PEATURE: PRT

ORGANISM: Artificial Sequence

SEQ ID NO 20

LENGTH: 535

OUTHER INFORMATION: Pfam consensus sequence

US-09-795-693-20

Ouery Match
Best Local Similarity 26.4%; Score 90.5; DB 10; Length 535;

Best Local Similarity 26.4%; Pred. NO. 0.27;

Best Local Similarity 26.4%; Pred. NO. 0.2
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Maximum DB seq length: 2000000000
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                                738.6
208.4
208.8
165.2
1165.2
110.2
90.2
80.6
62.6
62.6
59.4
9.8
90.2
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(without alignments) updates/sec
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Published_Applications_NA:*
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'cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
'cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
'cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
'cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
'cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
'cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
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'cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
'cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
'cgn2_6/ptodata/2/pubpna/US10_PUBCOMB.seq:*
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1 US-09-960-352-3386

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1 US-09-883-381-2007

1 US-09-917-800A-1646

1 US-09-917-800A-1646

1 US-09-917-807-1646

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1 US-09-917-918-743A-4

1 US-09-90-918-743A-2

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1 US-09-90-988-1

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   US-09-864-761-12350
US-09-969-373-354
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US-09-816-028A-30
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                                                                                                                                                                                         Sequence 4, Appli
Sequence 6, Appli
Sequence 3386, Ap
Sequence 8860, Ap
Sequence 25526, Ap
Sequence 2007, Ap
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Sequence 2, Appli
Sequence 4582, Ap
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                Sequence 354,
                              Sequence 30,
Sequence 123:
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| sequence 5558, Ap sequence 981, App sequence 17, Appl sequence 17, Appl sequence 10, Appl sequence 10, Appl sequence 17, Appl sequence 381, Appl sequence 376, Appl sequence 13489, A sequence 1356, Ap sequence 14521, A sequence 14521, A sequence 1250, Ap sequence 1250, Ap sequence 1250, Ap sequence 1200, Ap sequence 1200, A sequence 1200, A sequence 1200, Ap sequence 1200, Ap sequence 1200, Ap sequence 2100, Ap sequence 2100, Ap sequence 2100, Ap sequence 2100, Ap sequence 4201, Appl sequence 15, Appl |

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US-10-050-786-3
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APPLICANT: Ruben et al.
APPLICANT: Ruben et al.
TITLE OF INVENTION: Calcium Channel Polynucleotides, Polypeptides, and Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: FIGURER: US/10/050,786
CURRENT APPLICATION NUMBER: US/07/74,028
CURRENT EILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 09/774,028
PRIOR APPLICATION NUMBER: PCT/US00/20392
PRIOR PFLING DATE: 2001-07-27
PRIOR PFLING DATE: 1090-07-27
PRIOR PFLICATION NUMBER: US 60/145,958
PRIOR FILING DATE: 1990-07-28
PRIOR FILING DATE: 1990-07-28
PRIOR APPLICATION NUMBER: US 60/149,446
PRIOR APPLICATION NUMBER: US 60/149,446
PRIOR PILING DATE: 1990-08-18
PRIOR FILING DATE: 1990-08-18
PRIOR FILING DATE: 1900-03-14
PRIOR FILING DATE: 2000-03-14
PRIOR FILING DATE: 1900-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence 3, Application US/10050786 Patent No. US20020155539A1
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                                                                                                                                                                                                                  43.3%; Score 738.6; DB 9; Query Match
Best Local Similarity 98.7%; Pred. No. 1.9e-146; Best Local Similarity 0; Mismatches 9; Matches 755; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                ENGTH: 1316
                                                                                                                                                                                                                                                                                                                                                                                             ID NO 3
1053 CACCAGAAGGGTTTTTTTCCCTACAACCAGCAAGAACATATATTAGATACATGAATCTCA 1112
                                                                                                                                           933 AAAAAAAAATCTCAAGGAAAAGGCATAAAAATGATTGCTACACAAAAGTGACCAAATTTTTA 992
                                                              PatentIn Ver. 3.1
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1113 AȚȚAȚAAȚȚAȚGGCAȚȚAAȚȚȚGCAȚTȚTAȚȚTCAAAAȚȚAACȚTGGGGGGCACAȚGȚAAȚ 1172
                                                                       1173 ÇIÇTTGAÇÇAATÇTGATATTTTTGGGAAGTCÇTTTAAAAAGTTAÇAAATTTATÇAATAAA 1232
                                                                                                                        1233 TTACTAGTAGATAAGATGATTCAGAAACAAAAGAAATCACAGAATTAGGATGTGGCTGG 1292
646 CACCAGAAGGGTTTTTTCCCTACAACCAGCAAGAACATATATAGATACATGAATCTCA 705
                                      1293 CTGGTGTATGAAGCACCATGTGATGAATTCATAAAGTTGCAAAAGTCAAAACAAAAAATACTGT 1352
                                                                                        1353 ACATGCAACCAGAAATCAAAATAAATTAGAGACCTATATAAATGCATTTAATA 1412
                                                                                                                                                                                                                                         1413 CATGATACTTTTGACATAATAAGCCATTGGAAAAGGGAAAGATTAGATACTAAATAAGAT 1472
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                                                                                                                                                                                                                                                                                                                         TICTCTCTGTATATTTAAACAAATAGCTGGTATA 1592
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Sequence 4, Application US/10095139
Sequence 4, Application US/10095139
Sequence 4, Application US/2020165357A1
Septem Vo. US20020165357A1
APPLICANT: Curtis, Rory A.J.
APPLICANT: Silos-Santiago, Inmaculada
TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic
TITLE OF INVENTION: 10 Transporters and Uses Therefor
CURRENT APPLICATION NUMBER: US/10/095,139
FRIOR APPLICATION NUMBER: 60/275,172
NUMBER OF SEQ 1D NOS: 24
SOPTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1653 TÇAAAÇTÇTATÇTAÇATTTAATAAATAAÇTATAAATTAAAAAT 1697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
i LOCATION: (1562)
i O'THER INFORMATION: "m" represents ambiguous nucleotides
i LOCATION: (1)...(2866)
i OTHER INFORMATION: (1)...(2866)
i OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                          1533 TTTAATTACTTTTTCTTTGTAATTT
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SEQ ID NO 4
LENGTH: 2866
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155 GCCTCCTCTTGGGGGTCACGTGTTCACAATGGCCTTTGAGGAGGTCTTGAGTGAAG 214
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                                                                                                                                                   393 TGGGTGGCCTGGGCAGGTTCCAGGTTCTCCAGACGATGGCTCTGATGGTCTCCCATCATGT 452
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                                                                                                                                                                                                                                                                                                                                                                                455 GCTTTGTCCATCCCCAGTGGCAGCTTCTTCACCTGAATGGGACTATCCACAGGACAAGTG 514
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Query Match
Best Local Similarity 62.8%; Score 218.4; DB 9; Length 2866;
Matches 339; Conservative 0; Mismatches 201; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Curtis, Rory A.J.
APPLICANT: Glios-Santisgo, Inmaculada
APPLICANT: Milennium Pharmaceuticals, Inc.
TITLE OF INVENTION: 38554, 57301, and 5824, Human Organic
TITLE OF INVENTION: Ion Transporters and Uses Therefor
FILE REFERENCE: MPIOL-017PLRNM
CURRENT APPLICATION NUMBER: US/10/095,139
FRIOR APPLICATION NUMBER: 602-03-11
FRIOR PLING DATE: 2001-03-12
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FastSEQ for Windows Version 4.0
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OTHER INFORMATION: n represents ambiguous nucleotides
LOCATION: fac feature
LOCATION: (1662)
OTHER INFORMATION: n = A,T,C or G
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US-10-095-139-6
Sequence 6, Application US/10095139
Fatent No. US20020165357A1
GENERAL INFORMATION:
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ORGANISM: homo sapiens
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Best Local Similarity
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                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Bos taurus
OTHER INFORMATION: Clone ID: 15-LIB3058-028-Q1-K1-D3
US-09-960-352-3386
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                                                                                                                                                                                                                                                                                                                          SEQ ID NO 3386
LENGTH: 265
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APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
TITLE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
                                                                                                                                                            Matches 200; Conservative
                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Warren, Wesley C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       187 АГСССТТТСАССАССТСТТСАСТСААСТТССАСССТТСССАСАСТТССАСТССАТ 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         247 CTGGTTTTTATTCTTCCCTCTCATGTTATTAATCCCTCATATACTGCTAGAGAACTTT 306
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                                                                                                        258 TCTTCCCTCTCATGTTATTAATCCCTCATATACTGCTAGAGAACTTTGCTGCAGCCAT 317
    204
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TÉCTGGTCATCACTGCTGGGTCCACAATCTTGATAATGCCACTGTCTCTGATAATGATAC 145
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                                                                                TTTTCCTTTTTACATGATAGTAGTCTGTCACTCATTGCTGGAGAACTTCACTGCAGCCGT 205
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77.5%;
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                                                                                                                                                              Score 165.2; DB 10; Length 265; Pred. No. 4.6e-26; O; Mismatches 58; Indels 0;
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US-09-864-761-8860/c
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-02-04
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                                                                                                                                                                                                                                                                                                             SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 8860
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/006669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR APPLICATION NUMBER: PCT/US01/00668
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NUMBER OF SEQ ID NOS: 49117
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OTHER INFORMATION: MAP TO AC012153.10
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 6
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 7.2
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                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                             FEATURE:
                                                                                                                                                                                  ORGANISM: Homo sapiens
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Hanzel, David K.
Chen, Wensheng
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264 CTCTCTCATGTTATTAATCCCTCATATACTGCTAGAACTTTTGCTGCAGCATTCCTGG 323
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63.7%; Pred. No. 6.3e-20;
tive 0; Mismatches 118; Indels 0;
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OTHER INFORMATION: EXPRESSED IN BALIN, SIGNAL = 6
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 7.2
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 7
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 6.7
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN PETAL INTER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN PETAL INTER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL INTER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETALUTE 6.00e-79
OTHER INFORMATION: SWISSPROT HIT: Q920E8, EVALUE 3.00e-05
OTHER INFORMATION: EXT_HUMAN HIT: BF196052.1, EVALUE 1.00e-89
                       PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00601
PRIOR APPLICATION NUMBER: PCT/USO1/00601
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR PILING DATE: 2000-09-21
PRIOR PLING DATE: 2000-06-30
PRIOR PLING DATE: 2000-06-30
PRIOR PILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers: 1.1
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; Patent No. US20020132090A1
; GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Rank, David R.
APPLICANT: Hand, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
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                                                                                                                                                                                                                                                                                                                                                                                184 TCAATGGCCTTTGAGGAGCTCTTGAGTCAAGTTGGAGGCCTTGGGAAGTTTCAGATGCTT 243
                                                                                                                                                                                                                                                                                                                                                                                                                                  536 TCCATGGCATTTTCTGAACTCCTGGACCTCGTGGTGGCTGGGCAGGTTCCAGGTTCTC 477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           244 CATCTGGTTTTTATTCTTCCCTCTCATGTTATTAATCCCTCATATACTGCTAGAAC 303
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', OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 5.7
') OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 7
') OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5
') OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 6.1
') OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 6.9
') OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 6.9
US-09-864-761-8860
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Patent No. US20020048763A1
GENERAL INFORMATION:
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Best Local Similarity 62.5
Matches 253, Conservative
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US-09-917-800A-1646
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 3.0
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PRIOR FILING DATE: 2000-07-31
PRIOR RPPLICATION NUMBER: US 60/222,880
PRIOR FILING DATE: 2000-11-02
PRIOR APPLICATION NUMBER: US 60/290,029
PRIOR APPLICATION NUMBER: US 60/290,029
PRIOR FILING DATE: 2001-05-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Elashoff, Michael APPLICANT: Gene Logic, Inc.
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 401
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TITLE OF INVENTION: Molecular Toxicology Modeling
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                                       OR APPLICATION NUMBER: US 60/290,645
OR FILING DATE: 2001-05-15
OR APPLICATION NUMBER: US 60/292,336
OR APPLICATION NUMBER: US 60/292,336
OR FILING DATE: 2001-05-22
OR APPLICATION NUMBER: US 60/295,798
OR APPLICATION NUMBER: US 60/297,457
OR APPLICATION NUMBER: US 60/297,457
OR APPLICATION NUMBER: US 60/297,457
OR APPLICATION NUMBER: US 60/298,884
OR APPLICATION NUMBER: US 60/298,884
OR FILING DATE: 2001-06-19
OR FILING DATE: 2001-06-19
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FILING DATE: 2001-07-09
                        APPLICATION NUMBER: US 60/303,459
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Castle, Arthur
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                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4511, Application US/09983965
Patent No. US20020137160A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 268; Conservative
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                    SEQ ID NO 4511
LENGTH: 370
                                                                                APPLICANT: Byatt, John C.

APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 37-21(10297)C
CURRENT APPLICATION NUMBER: US/09/983,965
CURRENT FILING DATE: 2001-10-26
CURRENT FILING DATE: 1998-12-15
PRIOR FILING DATE: 1998-12-15
PRIOR APPLICATION NUMBER: US 60/113,678
PRIOR APPLICATION NUMBER: US 60/113,678
PRIOR FILING DATE: 1998-12-17
PRIOR FILING DATE: 1998-12-17
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                                                                   NUMBER OF SEQ ID NOS: 5912
TYPE: DNA
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Local Similarity 52.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGTCTATGACAACAGCACCTTCCCCTTCAACCATCGTGACTGAGTGGAACCTTGTGTGCT 638
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Pred. No. 5.5e-10;
0; Mismatches 208;
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DEPLICANT: Warren, Wesley C.
APPLICANT: Warren, Wesley C.
APPLICANT: Byatt, John C.
APPLICANT: Byatt, John C.
APPLICANT: Byatt, John C.
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND TITLE OF INVENTION: MUSCLE AND FAI DEPOSITION
TITLE OF INVENTION: WUMBER AND FAI DEPOSITION
TITLE OF INVENTION: WUMBER OF 37-21 (10298) C
CURRENT APPLICATION NUMBER: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 4582
LENGTH: 393
                                                                                                                                                                                                                                          APPLICANT: Neau, Jun-ichi
APPLICANT: Neau, Jun-ichi
TITLE OF INVENTION: SYSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
FILE REPERENCE: 06501-073001
CURRENT APPLICATION NUMBER: US/09/798,743A
CURRENT FILING DATE: 1909-09-07
PRIOR FILING DATE: 1999-09-07
PRIOR FILING DATE: 1999-09-07
PRIOR FILING DATE: 1998-09-07
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 1831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.5%; Score 59.4; DB 10; Length 1831; Best Local Similarity 55.6%; Pred. No. 0.0016; Matches 114; Conservative 0; Mismatches 91; Indels 0;
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OTHER INFORMATION: Clone ID: 20-LIB3057-013-Q1-K1-E7

US-09-960-352-4582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 4582, Application US/09960352; Patent No. US20020137139A1
                        570 AATGTGCTGTTTTTGACCATGGGCA 594
                                                                                                                                                                         ; Sequence 2, Application US/09798743A; Patent No. US20020099093A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY: CDS
; LOCATION: (124)..(1794)
US-09-798-743A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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CURRENT APPLICATION NUMBER: US/09/798,743A
CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: PCT/JP99/04853
PRIOR APPLICATION NUMBER: D10-252683
PRIOR FILING DATE: 1999-09-07
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                  124 chardescrincerrendedacherrendederrirangeschergenaraerrendaharenge 183
                                                                                                                                                                                                                                                                                                                                     245 AICTGGTTTTTATTCTTCCCCTCTCATGTTATTAATCCCTCATATACTGCTAGAACT 304
                                                                                                                                                                                                                                                                                                                                                                                   184 AGAİĞGAİİCACİİCİİÇİİÇİRİÇAAĞCİCAİACİCIGİCGGGİGGGAIİGĞIIAAÇİ 243
                                                                                                                                                                                                                                                                                                                                                                                                                                         TIGCIGCAGCCATICCTGGICATCGTTGCTGGGTCCACATGCTGGACAATAATACTGGAT 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         244 TCAGCGATGTGATACATCATCAACGTTGCAGGTCTCCATACTGGATTTGCACTGCCT 303
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                                                                                                                                           Length 370;
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Best Local Similarity 56.6%; Pred. No. 0.00034;
                                                                                                                           Query Match 4.7%; Score 80.6; DB 10; Length Best Local Similarity 57.9%; Pred. No. 3.2e-08; Matches 143; Conservative 0; Mismatches 104; Indels
                                                     ; OTHER INFORMATION: Clone ID: 08-LIB34-012-Q1-E1-B7
US-09-983-965-4511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/09798743A Patent No. US2002099993A1 GENERAL INFORMATION:
APPLICANT: Neau, Jun-ichi
APPLICANT: Ose, Asuka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       706 TCTGCTCGGTGGTTGATAATCACCA 730
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; LOCATION: (60)..(1730)
US-09-798-743A-4
ORGANISM: Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                425 ACTCAAA 431
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LENGTH: 1888
TYPE: DNA
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                                                                                                                     Db 527344 TGTTTTTTGATCTAAĀĀĀTAATTTTTTTGATAACĀTTTĞATTTATTCATAGTCAAAATG 527285
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US-09-790-988-1/c
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US-09-790-988-1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 640681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09790988 Patent No. US20020127687A1
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                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: JP2000-107160
PRIOR FILING DATE: 2000-04-07
NUMBER OF SEC. TO TO THE PRIOR FILING DATE: 2000-04-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SHIGENOBU, SHUJI
APPLICANT: WATANABE, HIDEMI
APPLICANT: WATANABE, HIDEMI
APPLICANT: HATTORI, MASAHIRA
APPLICANT: SAKAKI, YOSHIYUKI
TITLE OF INVENTION: GENOME DNA OF BACTERIAL SYMBIONT OF APHIDS
FILE REFERENCE: 081356/0159
CURRENT APPLICATION NUMBER: US/09/790,988
CURRENT FILING DATE: 2001-02-23
                                                                                                                                                                                                                                   1088 ACATATATTAGATACATGAATCTCAATTATAATTATGGCATTAATTTTGCATTTTATTTCA 1147
                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                              1148 AAATTAACTTGTGGGGACATGTAATCTCTTGAGCAATCTGATATTTTTTGGGAAGTCCTTT 1207
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                                                                               1208 АААААGTTACAAATTTATCAATAAATTACTAGTAGATAAGATGATTCAGAAACAAA-AGA 1266
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1267 AAATCACAGAATTAGGATGTGGCTGGCTGGTGTATGAAGCACCATGTGATGAATTCATAA 1326
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                                                                                                                                                                                                                                                                                  2.9%; Score 50; DB 10; Length 64(ilarity 44.7%; Pred. No. 1; Conservative 0; Mismatches 340; Indels
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45.9%;
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Db 526984 TTTTTTTAAATATGTTTTTCTCTTTTTTTTAATATTTTTTTAATTTTTTAATTTTCAGTTTTTATATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Db 526864 TTÁTTTTÁÁÁTTATÁGAAATÁAATTTTTTTÁÁÁÁTAGTÁAACÁATGTTATTÁÁÁTTAGATA 526805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Db 526924 TATGTÁTCATGAAAAAATAGATCTACTTTTTTTGAGTATGAATAAAATGTATTTTTCTTA 526865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 253; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.9%;
Best Local Similarity 45.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/790,988
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: JP2000-107160
PRIOR FILING DATE: 2000-04-07
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SHIGENOBU, SHUJI
APPLICANT: WATANABE, HIDEMI
APPLICANT: HATTORI, MASAHIRA
APPLICANT: SAKAKI, YOSHIYUKI
TITLE OF INVENTION: GENOME DNA OF BACTERIAL SYMBIONT OF
FILE REFERENCE: 081356/0159
                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Buchnera
                                                                                                                                                                                                         77959 ATAAGATCGCATCAAATA---ATTATTCACAACGAAATAGAAAAAGAGCAATATTCAAGAA 78015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 640681
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1202 TCCTTTAAAAAGTTACAAATTTATCAATAAATTACTAGTAGATAAGATGATTACGAAAACA 1261
                                                                                                                                                1142 ATTTCAAAATTAACTTGTGGGGACATGTAATCTCTTGAGCAATCTGATATTTTTGGGAAG 1201
                                                                                                                                                                                                                                                                1082 GCAAGAACATATATTAGATACATGAATCTCAATTATAATTATGGCATTAATTTGCATTTT 1141
                                                                                                                                                                                                                                                                                                                                                                                  1022 AATTCAGAAAAAAAATACAGGAAAAGAACACACCAGAAGGGTTTTTTTCCCTACAACCA 1081
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                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches 297; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 49.8; DI
Pred. No. 1.1;
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Sequence 3, Application US/09801861
Patent No. US20020119544A1
Patent No. US20020119544A1
APPLICANT: VAM, Chunhua et al.
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLO01098
CURRENT PAPLICATION NUMBER: US/09/801,861
CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 53332
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78076 AATAGTAAAAATTTAAAAAATAATTATATTCCTGTTGATAGTAAAATTAATCGTGATAAC 78135
                                                                                              78136 ACAGTAAATTATGAATTAAATCATAGTGTATCACATACAAAAATGAACATAGGAGAAATT 78195
                                                                                                                                                                                                                                                                                                 78376 AAAATACCAGTTCATATAAATCATATAAATACTTTTAGAAAATCAAATTTTTTATATAAT 78435
                                                                                                                                                                                              78196 AAAAGGTTATCAGCAGCAGTTATAGTGAACTTTTCTAAAGATAAAATGGAAAATTGTC 78255
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                                                                                                                                               1322 CATAAAGTTGCAAAAGTCAAAACAAT--ACTGTACATGCAACCAGAAATCAAAATAAATC 1379
                                                                                                                                                                                                                                                  1380 CAGAAATAGAGACCTATATAAAATGCATTTAATACATGATACTTTTGACATAATAAGCCAT 1439
                                                                                                                                                                                                                                                                                                                                                   1440 TGGAAAACGGAAAGATTAGATACTAAATAACATTGACTATCTCTTTGTAAATACAGTCAC 1499
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1500 TAAATGATGTTAGTTACTTTTCCATGGTGGAATTTTTAATTACTTTTTCTTTGTAATTTT 1559
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78436 Trrgcaccargerrr 78450
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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 53.2
49.6
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47.8
47.6
45.6
44.6
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1 aagaactgaggaagctcttt.....aaattaaaaaattatatttc 1705
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/2/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/2/ina/backfiles1.seq:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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2.58
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US-09-172-711-48
US-09-330-245A-1
US-09-572-147-1
US-08-232-463-14
US-08-647-397-1
US-08-647-397-1
US-08-647-397-1
US-08-98-416-541
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US-08-883-795A-36
US-08-107-755A-36
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US-08-98-416-535
US-08-98-416-332
US-08-998-416-332
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US-08-998-416-786
US-08-998-416-786
US-08-98-629-11
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                                  Sequence 48, Appl
Sequence 1, Appli
Sequence 1, Appli
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Sequence 1, Appli
Sequence 1, Appli
Sequence 5, Appli
Sequence 541, Appli
Sequence 542, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 36, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 8, Appli
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Sequence 8, Appli
Sequence 1137, Ap
Sequence 135, Appl
Sequence 37, Appl
Sequence 37, Appl
Sequence 37, Appl
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RESULT 2 US-09-330-245A-1

Sequence 1, Application US/09330245A Patent No. 6433631
GENERAL INFORMATION:
APPLICANT: GILEAD SCIENCES, INC. et

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| 305 TIGCT | 245 ATCTG 169 AGATG | 185 CAATG 109 CAATG | 26 49 | / Match Local Simi nes 161; | 1 2 8 P | 0 4 2 5 7 7 | ANT | 711- | | 41.8 41.8 | 42.2 42.3 | ,,,,, | 2444 | 2001 | υN |
| GCAGCC | ATCTGGTTTTTATTC AGATGGTTTTTGTTG | GGCCTTT | CTTTGAACTTA CTTTGTACTAA | milarity Conserva | ERL Pro | ENTION: ENTION: CATION CATION O ID NO: | ORMATION: Cunnigham, Zweiger, Ga | .48 Applica | | | | | , , , , , , , , , , | | |
| ATTCCTO | ATTCTTO | CAATGGCCTTTGAGGAGCTCTTGAGTC | 1-1 | 5. ty 70. ervative | ram 70 | II: SELLIAMEL, VELLEY, VILLEY, | m, Mary Gary B Scott R | t: | | 1519 9636 9636 | 6243 168575 782 | 1422 2334 837 | 732 828 834 | 701 724 | 19124 |
| 610 | TOTI HOTI | id—id | AAA - | * * | 771: | 10 71 | я. В. | 3/09 | | 411 | 044 | 4444 | 444 | 444 | 0 4 |
| TGCTGCAGCCATTCCTGGTCATCGTTGCTGGGTCCACATGCTGGACAAT ACTGCAGCCATTCCCAGTCATCGCTGCTGGGTTCCTATCCTCGACAAT | TCTGGTTTTATTCTTCCCTCTCTCATGTTATTAATCCCTCATATACTGCTAGAGAAC | CTTTGAGGAGCTCTTGAGTCAAGTTGGAGGCCTTGGGAGA | TCCGTAAAGCCATT-GTGCCTCCTCTTGGGGGTCACGTGTT | ; Score 98.8; DB 4; ; Pred. No. 3.8e-16; 0; Mismatches 67; | 146 | NG TOXICOLOGICAL RESPONSE US/09/172,711 | - W | on US/09172711 | ALIGNMENTS | 07-971-75 08-323-17 08-954-44 | US-09-056-075-1 US-09-426-290-1 US-09-007-119-15 | US-08-319-704-5 US-08-319-704-5 US-08-062-632-4 US-08-998-416-288 | 1 1 1 1 | -08-998-416- -08-998-416- -08-998-416- | 87-826E 98-416- |
| GGACAAT 354 CGACAAT 277 | TATACTGCTAGAGAACT 304 TATTATAATAGAGAAC- 227 | AAGTTGGAGGCCTTGGGAGATTTCAGATGCTTC 244 | CACAAT | Length 280; Indels 2; Gaps | | (S E S | | | | 1, | 15, | Sequence 5, Appli Sequence 4, Appli Sequence 288, App | Sequence 1036, App Sequence 538, App Sequence 305, App Sequence 2, Appli | 93 70 | Sequence 13, Appl Sequence 185, App |

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APPLICANT: John Feild
APPLICANT: Harma Ellot
APPLICANT: Harma Ellot
APPLICANT: HARMA ELLOT
TITLE OF INVENTION: POLYNUCLEOTIDE AND POLYPEPTIDE SEQUENCES
TITLE OF INVENTION: ENCODING MURINE ORGANIC ANION TRANSPORTER 5 (MOATPS) AND
TITLE OF INVENTION: SCREENING METHODS THEREOP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ñ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       185 CAATGGCCTTTGAGGAGCTCTTGAGTTCGAGGCCTTGGGAGATTTCAGATGCTTC 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.9%; Score 83.8; DB 4; Length 2123;
51.9%; Pred. No. 5.7e-12;
tive 0; Mismatches 212; Indels 33; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 365 CIGGIAATGAAACTGGAATCCTCAGTGAAGATGCCCTCTTGAGAATCTCTATCCCACTAG 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE: OTHER INFORMATION: Description of Unknown Organism:This information OTHER INFORMATION: is not available.
TITLE OF INVENTION: NOVEL GENE ENCODING ORGANIC ANION TRANSPORTER CURRENT APPLICATION NUMBER: US/09/330,245A
CURRENT FILING NOTE: 1999-06-10
PRIOR APPLICATION NUMBER: 60/088,864
PRIOR PILING DATE: 1999-06-11
PRIOR PILING DATE: 1999-06-11
PRIOR FILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 1
LENGTH: 2123
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US-09-572-147-1
; Sequence 1, Application US/09572147
; Patent No. 6420544
; GENERAL INFORMATION:
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Matches 264; Conservative
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LOCATION: (263)..(1912)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-330-245A-1
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517 GCAGACAGAACCCTGTGTGGATGGCTGGGTATATGATCAAAGCTACTTCCCTTCGACC 576
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  577 ATTGTGACTAAGTGGGACCTGGTATGTGATTATCAGTCACTGAAATCAGTGGTTCAATTC 636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          340 Arcárdachdadadchrdrardrachchrdadachtraceccaechderdada
                                                                                                                                                                                                                                                                                                                                               0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.8%; Score 81; DB 4; Length 1638; 66.1%; Pred. No. 2.7e-11; Live 0; Mismatches 60; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14, Application US/08232463
Patent No. 5670367
GENERAL INFORMATION:
APPLICANT: BORNER, F.
APPLICANT: FALKNER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSER: Foley & Lardner
STREET: 1800 Diagonal Road, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/232,463
FILE REFERENCE: GP-70622
CURRENT APPLICATION NUMBER: US/09/572,147
CURRENT FILING DATE: 2000-05-17
PRIOR PAPLICATION NUMBER: 60/134,879
PROR FILING DATE: 1999-05-19
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FARLSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 30472/114 IMMU
TELECOMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
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ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29.768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/935,313
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 22213-0299
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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nucleic acid
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                                                                                                                                                                                                    TYPE: DNA ORGANISM: MUS MUSCULUS
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                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                        SEQ ID NO 1
LENGTH: 1638
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     652 ATGCTGGTGGGAGGCATCATAGGTGGCCATGTCTCAGACAGGTGGCTGGTGGAATCTGCT 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1072 ССТАСЛАССАССАЛБАЛСАТАТАТТАБАТАСАТБААТСТСАА 1113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     772 CGCACAAATGGAATAAAGAATGCTGAAGAAACCCTGAACATAGAGGTTGTAAGATCCACC 831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  712 CGGTGGTTGATAATCACCAATAAACTAGATGAGGGCTTAAAGGCACTTAGAAAAGTTGCA 771
                                                                                                                                                                                                                                                                                                                      APPLICANT: Slade, Martin B
APPLICANT: Chang, Andy C M
APPLICANT: Williams, Keith L
APPLICANT: Williams, Keith L
TITLE OF INVENTION: Improved Plasmid Vectors for Cellular
TITLE OF INVENTION: Slime Moulds of the Genus Dictyostelium
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1049
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       952 AGGCATAAAAATGATTGCTACACAAAAGTGACCAAATTTTAAGAAGCCTTCATGAGCTGA 1011
                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5389526ris
STREET: One Liberty Place 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 3.5%; Score 60.4; DB 1; Length 7218; Local Similarity 8.7%; Pred. No. 8.4e-06; les 40; Conservative 228; Mismatches 194; Indels 0;
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          PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PJ 7187
APPLICATION NUMBER: PCT/AU90/00530
                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                      CITY: Philadelphia
                                                               APPLICATION NUMBER: US
FILING DATE: 19920625
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGCAGCCAAGCTCGGAATTAATTCTGTGAGCGTATGGCAA 1008
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                                                                                                                                                                                                                         USA
                                                                                                                                                                      Floppy disk
 02-NOV-1989
                                                                                     US/07/867,106
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ZIP: 02210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy

02210

COUNTRY: STATE:

USA

STREET:

600 Atlantic Avenue

Boston

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MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

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US-07-867-106-2
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INFORMATION FOR SEQ ID NO: 2:
                                                                                                    Sequence 1, Application US/08647397 Patent No. 5972702 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1405 ATTTAATACATGATACTTTTGACATAATAAGCCATTGGAAAACGGAAAGATTAGATACTA 1464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1345 АДТАСТОТАСАТОСААССАБАЛАТСАЛАТАЛАТСАБЛАДТАБАДССТАТАТАЛАТОС 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5491 TTTTAAGATTTTTTAATCTCGTCAATGATTTTAAAATAAAATCGATACATAATTTTAAA 5550
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1645 АССАДТІТІСЬВАСТСТАТСТАСАТТІВАТАВАТАВСТАТВВАТТЬВВАВ 1696
                                                                                                                                                                                                                                                                                                       1585 CTGGTATAGTTTACAATATTATAAAGATATTGTTCAAATTGAAGGGCCAAAGGCCAGGTTC 1644
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Feeney, Joanne Longo REGISTRATION NUMBER: 35,134
                         APPLICANT: Beier, David R.
APPLICANT: Brady, Kevin P.
TITLE OF INVENTION: OSTEOCLAST TRANSPORTER
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wolf, Greenfield & Sacks, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
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2378..5038
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,926B
                                      OPERATING SYSTEM: PC-DOS/MS-DOS
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Sequence 5, Application US/08731722

Patent No. 5961971
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Matches 217; Conservative
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MOLECULE TYPE: CDNA
                                                                                                                                                                                      FILING DATE: 10
CLASSIFICATION:
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Best Local S
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APPLICANT: Chitnis, Chetan
APPLICANT: Chitnis, Louis H.
APPLICANT: Buller, Louis H.
APPLICANT: Su, Xin-zhaun
APPLICANT: Su, Xin-zhaun
APPLICANT: Bullers, Thomas E.
TITLE OF INVENTION: AND PLASMODIUM FALCIPARUM ERYTHROCYTE BINDING PROTEINS
NUMBER OF SEQUENCES: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      479 ITCTTCACCTGAATGGGACTATCCACAGCACAAGTGAGGCAGACACAGAAGCCTGTGTGG 538
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1: 620 Newport Center Drive 16th Floor
Newport Beach
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                                                                                                                                                                                                                                                                  B0801/7048
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/647,397
                                                                                                      ATTCRNEY FACENT INFORMATION:
NAME: Gates, Edward R.
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: B0801
TELECOMUNICATION INFORMATION:
TELEPHONE: 617-720-3500
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LBNGTH: 2102 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
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Patent No. 5993827
GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
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120..1733
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MEDIUM TYPE: Floppy of
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US-08-647-397-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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15541 AAAATTATATAAATATATAAATTTTGATAGAATAAAAATGAAAAAGATTATCAAAAAA 15600
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                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 49.6; DB 2; Length 19124;
43.8%; Pred. No. 0.007;
tive 0; Mismatches 279; Indels 0;
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by Pythium oligandrum
                                      CLASSIFICATION: 435
ATTORNEY/AGATION: 435
ATTORNEY/AGATI INFORMATION:
NAME: ISRAELSEN, Ned
REGISTRATION NUMBER: 29,655
REFERENCE/DOCKET NUMBER: NIH121.001CP1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 235-850
TELEPHONE: (619) 235-850
INFORMATION FOR SEQ ID NO: 13: SEQUENCE CHARACTERISTICS:
LENGTH: 19124 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
US/08/487,826B
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TITLE OF INVENTION: Biocont:
TITLE OF INVENTION: by Pyth:
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NUMBER OF SEQUENCES:

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Best Local Similarity
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 1186 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Whitlock, Ted W.
REGISTRATION UNMBER: 36,'
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1244 TAAGATGATTCAGAAACAAAAGAAATCACAGAATTAGGATGTGGCTGGTGTATGA 1303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1184 TCTGATATTTTTGGGAAGTCCTTTAAAAAGTTACAAATTTATCAATAAATTACTAGTAGA 1243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1124 GGCATTAATTTGCATTTTATTTCAAAATTAACTTGTGGGGACATGTAATCTCTTGAGCAA 1183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
1544 TITCITIGIAATITITCICICIGIATATITIAAACAAATAGCIGGIATAGTITACAATAT 1603
                                                                                                                          1484 TTGTAAATACAGTCAACTAAATGATGTTAGTTACTTTTCCATGGTGGAATTTTAATTACTT 1543
                                                                                                                                                                                                                                                                                                                    1364 GAAATCAAAATAAATCCAGAAATAGAGACCTATATAAATGCATTTAATACATGATACTTT 1423
                                                                                                                                                                                                                                                                                                                                                                                                                   1304 AGCACCATGTGATGAATTCATAAAGTTGCAAAAGTCAAAACAATACTGTACATGCAACCA 1363
                                                                                                                                                                                                                         1424 ТGACATAATAAGCCATTGGAAAACGGAAAGATTAGATACTAAATAACATTGACTATCTCT 1483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 162 TTTTATTAACTTAAATTAAAAAGCGAATAAAATTAAGAAAGCAATCATTAAACAGAATAA 221
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                                                                                                                                                                                                                                                                      458 AAAATGAAAATACTTCTAATAAGAATATATTTATTATTATAAAATATTGAAAAACTTTTATTT 517
                                                                                                                                                                                                                                                                                                                                                                     401 CATATTATGTAATTTATATCTTTTTAAA----GATATATAACTTTTAAAAAAAAATT 457
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                                                                          578 ATAAGAAATTCTTCAATAGAGAAAGGTTTTATTTTTTCAATAGAAAATTGTAATTCTTAT 637
                                                                                                                                                                        518 ATATTTTTAAATAATGATCTGGATTTÄTÄCÄGTÄGÄTCCTÄTAAATCTTTAATAGTAGTT 577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 1186 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/731,722
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCACCTGCTAATCCAATAGTAGCTAATCCAGCACCAATAAATTTTTGCAGATTGT-AATAA 400
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Patent No. 6239264
                                                                                                                                 Matches
                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 919-541-8689 INFORMATION FOR SEQ ID NO:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1664 GTACATTTAATAAAATAACTATAAATTAAAAAAT 1697
                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PF/5-30306/A/CGC1976
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: GENOMIC DNA SEQUITITLE OF INVENTION: AND USES THEREOF
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1505 GATGTTAGTTACTTTTCCATGGTGGAATTTTAATTACTTTTTTCTTTTGTAATTTTTTCTCTC 1564
                                                                                          MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 821 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               638 TCTATTTTAATAAATAATCGTAATACTTTAATAAGAAATA----
                                                      732 AAGGGTAANINTAAATGCAAATTATTATTAGGGGTCGTTTINTCATTTTTATTATTATIATNA 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: No. 6239264artis Corporation STREET: 3054 Cornwallis Road CITY: Research Triangle Park STATE: No. 6239264th Carolina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: CH 0016/97 FILING DATE: 31-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 2
CLASSIFICATION:
                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/998,416
                                                                                                                                                 Local
                                                                                                                                                                                                                           ORGANISM: PAG1378RP
                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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                                                                                                                                                    Similarity
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VENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Philippsen, Peter
Pohlmann, Rainer
Steiner, Sabine
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Knechtle, Philipp
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                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                                                                                  single
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                                                                                                                                 Score 47.8; DB 4; Length 821;
Pred. No. 0.0054;
0; Mismatches 136; Indels
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STATE: Ontario
                                                Toronto
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STRANDEDNESS:
      ADDRESSEE:
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                                                                                        COUNTRY:
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PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR RILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674

LENGTH: 1983
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                                       1565 IGTATATITTAAACAAATAGCIGGTATAGTITTACAATATTATAAAGATATTGTTCAAATT 1624
                                                                                                                          1301 TGAAGCACCATGTGATGAATTCATAAAGTTGCAAAAGTCAAAACAATACTGTACATGCAA 1360
                                                                                  1361 CCAGAAATCAAAATAAATCCAGAAATAGAGACCTATATAAATGCATTTAATACATGATAC 1420
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672 AATNANAATACTTATTATATTTTCATTAATAATATTAAATNTNTTAAAAATAATTATAA 613
                                                                                                                                                                  216 TGATGATAAAAAAGCACTTCTGACAGGACAAGGACAAAGACTCTAATAAACCATTATCAGC 275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.8%; Score 47.6; DB 4; Length 1983; Best Local Similarity 49.6%; Pred. No. 0.0088; Matches 122; Conservative 0; Mismatches 124; Indels 0;
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APPLICANT: Awang, Gregor
TITLE OF INVENTION: Recombinant DNA Molecules and Expression
TITLE OF INVENTION: Vectors for Tissue Plasminogen Activator
NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                        Sequence 1524, Application US/09134001C Patent No. 6380370 GENERAL INFORMATION:
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US-09-134-001C-1524
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Patent No. 5985607
GENERAL INFORMATION:
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US-09-134-001C-1524
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1367 ATCAAAATAAATCCAGAAATAGAGACCTATATAAATGCATTTAATACATGATACTTTTGA 1426
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APPLICANT: REYMOND, CHRISTOPHE DOMINIQUE
TITLE OF INVENTION: DICTYOSTELID EXPRESSION VECTOR AND
TITLE OF INVENTION: METHOD FOR EXPRESSING A DESIRED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.7%; Score 45.6; DB 2; Length 665; Best Local Similarity 50.0%; Pred. No. 0.018; Matches 170; Conservative 0; Mismatches 164; Indels
                                                                                                            ZIP: MSH 312
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IPM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 322 TTTTATAATTAAATGTTTATAATTACATATTTATAATT 361
                                                                                                                                                                                                                                                     27-JUN-1997
27-JUN-1997
27-JUN-1997
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                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883
FILING DATE: 27-UUN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: GRAVALILE, MICHALINE
REFERENCE/DOCKET NUMBER: 7841
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 364-7311
TELEFAX: (416) 361-1398
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                           40 King Street West
BERESKIN & PARR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 665 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
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M5H 3Y2
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NUMBER OF SEQUENCES:

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                                                                                                     US-07-867-106-2/c
                                                                                                                       RESULT 13
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                                               Sequence 2, Application US/07867106
Patent No. 5389526
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2
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LENGTH: 731
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PRIOR APPLICATION NUMBER:
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MEDIUM TYPE: 3.5" FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1173 СТСТТGAGCAATCTGATATTTTTGGGAAGTCCTTTAAAAAGTTACAAATTTATCAATAAA 1232
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                                                                                                                                                                                                                                                                               1413 САТGАТАСТТТТGАСАТААТААGСССАТТGGAAAACGGAAAGATTAGATACTAAATAACAT 1472
                                                                                                                                                                                                                                                                                                                                                                                                                      1353 АСАТССААССАСАЛАЛТСАЛАЛТАЛАТССАСАЛАТАСАСАССТАТАТАЛАТССАТТТАЛТА 1412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             130 TTTCTATTTGAAGGAGTTTTTATTTGTATTTAAAATTATATTAAACATAGTGAACCTAAA 189
APPLICANT: Slade, Martin B
APPLICANT: Chang, Andy C M
APPLICANT: Williams, Keith L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: NUCLEIC ACID STRANDEDNESS: SINGI TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: Midwest Micro 486-50
OPERATING SYSTEM: DOS
SOFTWARE: WORDPERFECT 6.1
                                                                                                                                                                           430
                                                                                                                                                                                                                                            370 ATCTTTAATTGTGCAAAACACACTTTTAACACACTCTATTATCTTACAAAGGTTTAAAAT
                                                                                                                                                                                                                                                                                                                                                                                    CITY: PITTSBURGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 ATTTTTGTGTTCTTATAATTTGGTTAAATCGATGAATAATATTTGATTAGTATATGTTTT 69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 2.6%; Score 44.6; DB 1; Length 731; Local Similarity 43.7%; Pred. No. 0.034; es 197; Conservative 0; Mismatches 254; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTAATTACTTTTTCTTTGTAATTTTTCTCT 1563
                                                                                                                                                                                                                                                                                                                TTTAATTTTTTTTTATTTATTATTTTTTTTT 460
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Best Local Similarity 49.7%; Pred. No. 0.11;
Matches 196; Conservative 0; Mismatches
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 1992062
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                          1432 TAAGCCATTGGAAAACGGAAAGATTAGATACTAAATAACATTGACTATCTCTTTGTAAAT 1491
                                                                                                                                                                               1372 ААТАААТССАGАААТАGAGAССТАТАТАААТGСАТТТААТАСАТGATACTTTTGACATAA 1431
                                                                                                                                                                                                                 2259 АВАЙАТЖАЛАЛАЛАЙАЙАЛАЛАGКАЙАТТТТЖАЙАЙАЙАЛАЛАЙАЛАЙАЛТТАЙТСЛАЙА 2200
                                                                                                                                                                                                                                                      1312 GTGATGAATTCATAAAGTTGCAAAAGTCAAAACAATACTGTACATGCAACCAGAAATCAA 1371
                                                                                                                                                                                                                                                                                        2318 AT-TTACAATAAATAAATAAATAAAATAAAATGAAAGTGTGGGAAAAATAAAATTAAAAATT 2260
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1492 ACAGTCACTAAA--TGATGTTAGTTACTTTTCCATGGTGGAATTTTAATTACTTTTTCTT 1549
                                                                        2199 AATAAATATAATTAAAATTGTCATGCCAAAACTGATAAATATTTGATATATTATCCAATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Feeney, Joanne REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: AU PJ 7187
APPLICATION NUMBER: PCT/AU90/00530
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One Liberty Place 46th Floor
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2378..5038
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52 TATTTTAATTATAAAATATTTAA 25
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 904-375-8100
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INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy
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STRANDEDNESS: double
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ORIGINAL SOURCE:
ORGANISM: Amss
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LOCATION:
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NAME/KEY:
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US-07-991-867B-8
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                                                                                                      NEBOLI 1. TO SEQUENCE 36, Application US/08883795A
; Sequence 36, Application US/08883795A
; Patent No. 5985607
; GENERAL INFORMATION:
; APPLICANT: Delcuve, Genevieve
; APPLICANT: Delcuve, Genevieve
; TITLE OF INVENTION: Recombinant DNA Molecules and Expression
; TITLE OF INVENTION: Vectors for Tissue Plasminogen Activator
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: BERESKIN & PARR
; STREET: 40 King Street West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3;
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2.6%; Score 43.6; DB 2; Length 665;
Best Local Similarity 47.9%; Pred. No. 0.058;
Matches 186; Conservative 0; Mismatches 199; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                     2024 TTTTTTTTTTCAAGTAAAAAAAAAAAA 1991
1550 TGTAATTTTTCTCTGTATATTTTAAACAAATA 1583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Gravelle, Micheline
REGISTRATION NUMBER: 40,261
REFERENCE/DOCKET NUMBER: 7841-062
TELECOMMUNICATION INFORMATION:
TELEPHORE: (416) 364-7311
TELERAX: (416) 361-1398
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 665 base pairs
TYPE: nucleic acid
STRANDENDESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/883,795A
FILING DATE: 27-JUN-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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; IMMEDIATE SOURCE:
; CLONE: Rh 32
US-08-883-795A-36
                                                                                                                                                                                                                                                                                                                                                                                         E: Ontario
FRY: Canada
MSH 3Y2
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MOLECULE TYPE: c
ORIGINAL SOURCE:
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1614 TIGITCAAATIGAAGGGCAAAGGCCAGGTICAGCAATTITCAAACTGTATGTACATTTAA 1673
                                                       Sequence 8, Application US/07991867B
Patent No. 5476781
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL Hall, Richard W.
APPLICANT: Hall, Richard L.
APPLICANT: Gruidl, Michael E.
TITLE OF INVENTION:
OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: David R. Saliwanchik
STREET: 2421 N.W. 41st Street, Suite A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/991,867B
FILING DATE: 12-DEC-1992
CLASSIFICATION NUMBER: W0 92/14818
FILING DATE: 12-EEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: W0 07/827,685
FILING DATE: 30-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,685
FILING DATE: 19-EEB-1991
ATTORNEY/AGENT INFORMATION:
NAME: Saliwanchik, David R.
REGISSTRATION NUMBER: 31,794
PURDER SALIWANCHIN NUMBER: 31,794
PURDER SALIWANCHIN NUMBER: 31,794
PURDER SALIWANCHIN NUMBER: 31,794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1136 CATTTTATTTCAAAATTAACTTGTGGGGACATGTAATCTCTTGAGCAATCTGATATTTTT 1195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1076 СААССАВСААВААСАТАТАТТАВАТАСАТВААТСТСВАТТАТАТТАТВЕСАТТААТТВ 1135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1005 AATATTTTAAATAATATTCCAGAAAATATTAAAAGTTTATATATTTCAA------AT 1055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1196 GGGAAGTCCTTTAAAAAGTTACAAATTTATCAATAAATTACTAGTAGATAAGATGATTCA 1255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1316 ТСААТТСАТАААСТТССАЛААСТСАЛААСТАТАСТСТАСАТССААССАСАЛАТСАЛАЛТА 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1116 TCTTATAACAAAATAGCAATATAAGTAATATTATACTACCACATTCTATAGAATTTTTA 1175
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1376 AATCCAGAAATAGAGACCTATATAAATGCATTAATACATGATACTTTTGACATAATAAG 1435
                                                                               1292 TATAGTTGAGTTAAATATGGAATCAATACAAATAAAGATTATAAAATTTATAGAAAATT 1351
                                                                                                                                                                                                         1556 TITTCTCTCTGTATATTTTAAACAAATAGCTGGTATAGTTTTACAATATTATAAAGATATT 1615
                                                                                                                                                                                                                                                       1232 ARAAAAATTAATAATATCTAAAAATAAATTTGGTAACTTTAATAATGTTTTTCCTATTAG 1291
                                                                                                                                                                                                                                                                                  1496 TCACTAAATGATGTTAGTTACTTTTCCATGGGGGAATTTTAATTACTTTTTCTTTGTAAT 1555
                                                                                                                                                                                                                                                                                                                                         1176 AATTGTGAATCATGTAATATAAATGACTATA----ATTTTATTAATAATTTAGTAAATTT 1231
                                                                                                                                                                                                                                                                                                                                                                             1436 CCATTGGAAAACGGAAAGATTAGATAACTAACATTGACTATCTCTCTTTGTAAATACAG 1495
  1412 AAAATTTCCAAAAAGTATAA 1431
                                                                                                                         1616 GTTCAAATTGAAGGGCAAAG-GCCAGGTTCAGCAATTTTCAAACTGTATGTACATTTAAT 1674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               825 CAATGAAAAAAAAAATTATCAAAATTGGATTTACTAAATTCTGATATAATTTTTAATA 884
                                          1675 ААААТААСТАТАААТТАААА 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               945 ATTAATATATTAAAAAAATTAGTTAATTTAGAAGAATTGCATATAATATATTATGATAAT 1004
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Search completed: January 6, 2003, 22:45:58 Job time: 115.674 secs

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OM nucleic - nucleic search, using sw model
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0% Maximum Match 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Published Applications NA:*

1: /cgm2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgm2_6/ptodata/2/pubpna/US06_NEW PUB.seq:*

3: /cgm2_6/ptodata/2/pubpna/US06_NEW PUB.seq:*

4: /cgm2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgm2_6/ptodata/2/pubpna/US07_NEW PUB.seq:*

6: /cgm2_6/ptodata/2/pubpna/US08_NEW PUB.seq:*

7: /cgm2_6/ptodata/2/pubpna/US08_NEW PUB.seq:*

8: /cgm2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

9: /cgm2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

10: /cgm2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

11: /cgm2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

12: /cgm2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

13: /cgm2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

14: /cgm2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     381593 seqs, 216252194 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           January 6, 2003, 20:58:28; Search time 39.0945 Seconds (without alignments) 8894.701 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IDENTITY_NUC Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| 0 10 11 12 13 14 15 16 16 19 | c 2 7 6 | α . υ | C 4 | C C | N | _ : | Result No. |
|--|---|---------------------|--------------------|--------------------|-------------------|-------------------|-----------------------|
| 3 3 3 3 3 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 | 110.2 | 136.2 | 160.4 | 165.2 | 208.8 | 208.8 | Score |
| + + + + + + + + + + + + + + + + + + + | 13.7 | 16.9 | 20.0 | 20.5 | 26.0 | 26.0 | Query Match Length |
| 370 1888 1831 454 796 810 275 406 2189 302250 | 401 | 1339 | 540 | 265 | 2866 | 1662 | Length |
| 100 100 100 100 100 100 | 10 | ٥٢ | 10 | 10 | ø | 9 | DB |
| US-09-983-965-4511 US-09-798-743A-2 US-09-798-743A-2 US-09-764-887-47 US-09-764-887-376 US-09-974-300-945 US-09-974-300-945 US-09-9878-574-8151 US-09-950-352-11064 US-09-962-332-154 US-09-962-332-154 US-09-962-332-154 | US-09-833-381-2007 US-09-917-800A-1646 | US-09-864-761-20020 | US-09-864-761-8860 | US-09-960-352-3386 | US-10-095-139-4 | US-10-095-139-6 | ID |
| sequence 4,511, Appli Sequence 2, Appli Sequence 2, Appli Sequence 376, Appl Sequence 376, Appli Sequence 945, Appli Sequence 945, Appli Sequence 1164, Appli Sequence 154, Appli Sequence 456, Appli Sequence 45, Appli | Sequence 2007, Ap Sequence 1646, Ap | Sequence 3, Appli | sequence obov, as | Seguence Jose, Ap | riddy 't acmanbas | Sequence 6, Appli | Description |

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| 30 | 30.2 | 30.4 | 30.4 | 30.4 | 30.4 | 30.4 | 30.4 | 30.4 | 30.6 | 30.6 | 30.6 | 30.6 | 30.8 | 31 | 13 | 31 | 31.2 | 31.2 |) t | | י ני ני | 31.4 | 31.6 | 32 | 32 | 36.4 | J | |
| 3.7 | 3. 8 | 3.8 | 3.8 | ა | ω | 3.8 | 3.8 | 3. 8 | œ | | 3.8 | 3.8 | ω | <u>د</u> | | | | · | ٠, | , c | | | | | | | | |
| 278 | 305 | 3980 | 3820 | 1563 | 432 | 432 | 432 | 349 | 174493 | 2019 | 2000 | 406 | 1870 | 62804 | 26/6 | 1312 | 166/KT | 1000 | 2001 | 25.0 | 437 | 4858 | 4527 | 1135 | 4 4 0 | 1 6 | 725 | |
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| US-09-923-876-4018 | US-09-960-352-890 | 08-10-002-600-67 | US-09-904-040-0 | 0 0 | 0 0 | US-09-810-936-176 | 03-03-24-400-176 | US-09-853-386-46 | US-09+804-47EA-0 | US-09-801-566-199 | US-09-938-842A-4409 | S-102-282-282-27-400 | US-U9-938-846A-6730 | 12 00 000 00 00 00 00 00 00 00 00 00 00 0 | C2+03-030-0465-1010 | 0101-4578-959 | 00-00-026 240 0 | 116-09-822-246-3 | -897-99- | 50 | US-09-864-761-4863 | US-09-954-456-733 | -901-940-5 | י טע | י פע | 2 | US-09-910-943-70 | |
| Sequence soro, of | 200 | 90 | æ ' | | Seguence 4547, Ap | | Semience 176. App | ٠. ١ | Semience 48. Appl | Semience 3. Appli | | Seguence 4409. Ap | Semience 5140. Ap | Semience 2756, Ap | Sequence 3, Appli | semience 1919, Ap | 15 | Sequence 3, Appli | Sequence 5, Appli | Sequence 9965, Ap | Sequence 4600, Ap | | | Semience 3. Appli | | Seguence 967, App | 774 | |

ALIGNMENTS

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APPLICANT: Curtis, Rory A.J.

APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Milennium Pharmaceuticals, Inc.
APPLICANT: Milennium Pharmaceuticals, Inc.
APPLICANT: Milennium Pharmaceuticals, Inc.
APPLICANT: Milennium Pharmaceuticals, Inc.
TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic
TITLE OF INVENTION: 38554, 57301, and Uses Therefor
FILE REFERENCE: MPI01-017P1RNM
FILE REFERENCE: MPI01-017P1RNM
CURRENT FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: 60/275,172
PRIOR FILING DATE: 2001-03-12
NUMBER OF SEQ ID NOS: 24
NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 6, Application US/10095139 ; Patent No. US20020165357A1
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                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: n = A, T, C or G US-10-095-139-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 6
                                                                                                                                                                                                                                          Query Match 26.0%; Score 208.8; DB 9; Length 1662; Best Local Similarity 63.2%; Pred. No. 1.2e-53;
                                                                                                                                                                                                                      Matches 321;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: CDS
LOCATION: (1)...(1662)
OTHER INFORMATION: n represents ambiguous nucleotides
NAME/KEY: misc_feature
LOCATION: (1)...(1662)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 1662
TYPE: DNA
ORGANISM: homo sapiens
61 ACGATGGCTCTGATGGTCTCCATCATGTGGCTGTGTACCCAGAGCATGCTGGAGAAACTTC 120
                                                 61 CTGGTTTTTATTCTTCCCTCTCATGTTATTAATCCCTCATATACTGCTAGAGAACTTT 120
                                                                                                        1 Aragecarrirergaacrecragaceregrageregeciregecaegriceaegrierecae 60
                                                                                                                                                         1 ATGGCCTTTGAGGAGCTCTTGAGTCAAGTTGGAGGCCTTGGGAGATTTCAGATGCTTCAT 60
                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                            0; Mismatches 187;
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481 ATCATAGGTGGCCATGTCTCAGACAGGT 508
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121 GCTGCAGCCATTCCTGGTCATCGTTGGTGCTGGTGGTGGAGAATAATACTGGATCT 180
                                                   121 TCGGCCGCCGTGCCCAGCCAGCCGCTGCTGGGCCCCCTCCTGGACAGCACGCTCCG 180
                                                                                                                                     181 GGTAATGAAACTGGAATCCTCAGTGAAGATGCCCTCTTGAGAATCTCTATCCCACTAGAC 240
                                                                                                                                                                                                        181 GccAccarccradGGAGCricAdGiccrdAGGCCriccrGGCrAirriccAiccicGCGGGC 240
                                                                                                                                                                                                                                                                              241 TCAAATCTGAGGCCAGAGAAGTGTCGTCGCTTTGTCCATCCCCAGGGGAGCTTCTTCAC 300
                                                                                                                                                                                                                                                                                                                                      301 CIGAAIGGGACIAICCACAGAGACAGAGAGACACACAGAACCCIGIGIGGAIGGCIGG 360
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APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Milonnium Pharmaceuticals, Inc.
TITLE OF INVENTION: 38554, 57301, and 58324, Human Organic
TITLE OF INVENTION: 1on Transporters and Uses Therefor
FILE REPERENCE: MPIO1-017PIRNM
CURRENT APPLICATION NUMBER: US/10/095,139
PRIOR APPLICATION NUMBER: 60/275,172
PRIOR PELING DATE: 2001-03-11
PRIOR PILING DATE: 2001-03-12
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: (365)...(2026)
; OTHER INFORMATION: "n" represents ambiguous nucleotides
; NAME/KEY: misc feature
; LOCATION: (1)...(2866)
; OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                481 ATCATAGGTGGCCATGTCTCAGACAGGT 508
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Patent No. US20020165357A1
GENERAL INFORMATION:
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Sequence 3386, Application US/09960352

Patent No. US20020137139A1

GENERAL INFORMATION:

APPLICANT: Tao, Nengbing

APPLICANT: Byatt, John C.

TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION

TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION

TITLE OF INVENTION: WUSCLE AND FAT DEPOSITION

CURRENT APPLICANTE: 16511.006/37-21(10298)C

CURRENT APPLICANTON NUMBER: US/09/960,352

CURRENT FILING DATE: 2001-09-24

CURRENT FILING DATE: 2001-09-24
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485 TCGGCCGCCGTGCCCAGCCACCGCTGCTGGGCACCCCTCCTGGACAACAGCACGGCTCAG 544
                                                          181 GGTAATGAAACTGGAATCCTCAGTGAAGATGCCCTCTTGAGAATCTCTATCCCACTAGAC 240
                                                                                                                                                                             241 TCAAATCTGAGGCCAGAGAAGTGTCGTCGCTTTGTCCATCCCCCAGTGGCAGCTTCTTCAC 300
                                                                                                               545 dcchdcarccraddaddrigadtccrdaddccrtcradacratrrccardccardccadd
                                                                                                                                                                                                                     665 CCCAATGCCACGGCCACCAGCTGGAGCGAGCGAGCCGTGTGTGGATGGCTGG 724
                                                                                                                                                                                                                                                                                                                                                                                                               361 GTATATGATCAAAGCTACTTCCCTTCGACCATTGTGACTAAGTGGGACCTGGTATGTGAT 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 421 TATCAGTCACTGAAATCAGTGGTTCAATTCCTACTTCTGACTGGAATGCTGGTGGGAGGC 480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      785 İCİCATGCICİMGAĞCCCAİĞĞCCCAĞİCATCTACCİĞĞCİĞĞĞAİTCİĞĞTĞĞĞĞAĞCT 844
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                                                                                                                                                                                                                                                                                               301 CTGAATGGGACTATCCACAGCACAAGTGAGGCAGACACAGAACCCTGTGTGGATGGCTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Bos taurus
; OTHER INFORMATION: Clone ID: 15-LIB3058-028-Q1-K1-D3
US-09-960-352-3386
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US-09-864-761-8860/c
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                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Anno
                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00670 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: US 60/234.687
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CURRENT FILING DATE: 2001-05-23
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                                                                                                     OTHER INFORMATION: MAP TO ACO12153.10
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 6.2
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 7.2
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 5.7
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 7
OTHER INFORMATION: EXPRESSED IN LING, SIGNAL = 6.1
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 6.1
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.8
                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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                Local Similarity
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APPLICATION NUMBER: US 60/207,456
FILING DATE: 2000-05-26
APPLICATION NUMBER: US 09/632,366
FILING DATE: 2000-08-03
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/236,359
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APPLICATION NUMBER: US 09/608,408
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APPLICATION NUMBER: PCT/US01/00661
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Hanzel, David K.
Chen, Wensheng
Conservative
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  0
  Score 160.4; DB 10;
Pred. No. 3.8e-39;
0; Mismatches 151;
                                               Length
                                                    540;
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       Gaps
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PRIOR FILING DATE: 2000-10-04
PRIOR PELICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: Acomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
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APPLICANT: Rank, David F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00665

PRIOR APPLICATION NUMBER: PCT/US01/00668
                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
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                                                                        APPLICATION NUMBER: PCT/US01/00661
                                                                                                                        APPLICATION NUMBER: PCT/US01/00662
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
                                                                                                  FILING DATE:
                                                                                                                                                   FILING DATE: 2001-01-30
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Hanzel, David K.
                                                                                                     2001-01-30
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APPLICANT: Ruben et al.

TITLE OF INVENTION: Calcium Channel Polynucleotides, Polypeptides, and Antibodies
FILE REFERENCE: PT013PICI.

CURRENT APPLICATION NUMBER: US/10/050,786

CURRENT FILING DATE: 2002-01-18

PRIOR APPLICATION NUMBER: US/774,028

PRIOR APPLICATION NUMBER: PCT/US00/20392

PRIOR FILING DATE: 2000-07-27

PRIOR FILING DATE: 2000-07-27

PRIOR FILING DATE: 10999-07-28

PRIOR FILING DATE: US 60/145,958

PRIOR APPLICATION NUMBER: US 60/149,446
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OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 6
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 7.2
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 7.2
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5.7
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 5.7
OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 6.1
OTHER INFORMATION: EXPRESSED IN HEARY, SIGNAL = 6.1
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.9
OTHER INFORMATION: SWISSPROT HIT: G11441373, EVALUE 3.00e-05
OTHER INFORMATION: SWISSPROT HIT: BF196052.1, EVALUE 1.00e-89
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                                           PRIOR FILING DATE: 2000-09-21
PRIOR PLING DATE: 2000-09-21
PRIOR PPLICATION NUMBER: US 09/608,408
PRIOR PPLICATION NUMBER: US 09/774,203
PRIOR PILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers: 1.1
SEQ ID NO 25526
PRIOR APPLICATION NUMBER: US 60/234,687
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Patent No. US20020155539A1
GENERAL INFORMATION:
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Matches 207; Conservative
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ORGANISM: Homo sapiens
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Sequence 2007, Application US/09833381

Patent No. US20020132090A1

GENERAL INFORMATION:
APPLICANT: Robison, Keith E.

TITLE OF INVENTION: No. US20020132090A1e1 Nucleic Acid and Protein Homologs
FILE REFREENCE: 8000-119

CURRENT APPLICATION NUMBER: US/09/833,381

CURRENT FILING DATE: 2001-04-11

PRIOR FILING DATE: 2000-02-29

NUMBER OF SEQ ID NOS: 2050

SOFTWARE: FAILSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       679 ACTACTGTGTGTGACTTGTTCCGCAACCCCAGTATGCGTAAAAGGATCTGTATCCTGGTA 738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69 ACTACTGTGTGTGACTTGTTCCGCAACCCCAGTATGCGTAAAAGGATCTGTATCCTGGTA 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      315 CCACAGGACAAGTGAGGCAGACAC-AGAACCCTGTGGATGGCTGGGTATATGA-TCAA 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  373 AGCTACTICCCTICGACCATIGIGACTAAGIGGGACCTGGIAIGIGATIAICAGICACTG 432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                390 CCCCAACACAAATGAGCCAGACACGNNAGCCCTGTGTGGGTGGCTGGGTGTACGACAGAN 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    270 AAATCAANGGTTCAATCCCTATTTATGGCTGGGTCACTGCTGGGGGGGTCTAATAATGGC 211
                                                                                                                                                                                                                                                                                                                                                                                           9 CTGAACATAGAGGTTGTAAGATCCACCATGCAGGAGGAGGAGGAGGATGCAGCACAGAA
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Best Local Similarity 95.1%; Pred. No. 6.9e-30;
Matches 135; Conservative 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
13.7%; Score 110.2; DB 10; Length 401;
Best Local Similarity 73.9%; Pred. No. 6.9e-24;
Matches 164; Conservative 0; Mismatches 56; Indels 2;
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PRIOR FILING DATE: 1999-08-18
PRIOR APPLICATION NUMBER: US 60/189,064
PRIOR FILING DATE: 2000-03-14
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN Ver. 3.1
SEQ ID NO 3
LENGTH: 1316
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LOCATION: (1)...(401)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        739 TTTTGAGAAAAAATCTCAA 760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     129 TTTTTGAGATTTGCAAACACAA 150
                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-050-786-3
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Best Local Similarity
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US-09-833-381-2007/c
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LENGTH: 401
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RESULT 8

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US-09-917-800A-1646
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SEQ ID NO 1646
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Best Local Similarity 52.7%; Pred. No. 3.1e-17
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TITLE OF INVENTION: Molecular
FILE REFERENCE: 44921-5038-US
CURRENT APPLICATION NUMBER: US/09/917,800A
CURRENT FILING DATE: 2001-07-31
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PRIOR FILING DATE: 2000-07-31
PRIOR APPLICATION NUMBER: US 60/222,880
PRIOR FILING DATE: 2000-11-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 1740
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PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/298,884
PRIOR FILING DATE: 2001-06-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           314 GTCACCATGGTGGTTGCTCCCCTACTGCTGATGGCTTCCCACAACACCCTTGCAGAACTTC 373
361 GTATATGATCAAAGCTACCTTCCACCATTGTGACTAAGTGGGACCTGGTATGTGAT 420
                                                                           524 TACAATGGCAC---AGAAGCCAATGGCACCAGAGTCACAGAGCCCTGCATTGATGGCTGG 580
                                                                                                                                                                       464 AAGCAAGGACAACCCGAATCGTGCCTCCGCTTTACTTCCCCCCAGTGGGGGACCACCCTTT 523
                                                                                                                                                                                                               241 TCAAATCTGAGGCCAGAGAAGTGTCGTCGCTTTGTCCATCCCCAGTGGCAGCTTCTTCAC 300
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APPLICATION NUMBER: US 60/290,645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/295,798 FILING DATE: 2001-06-06
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APPLICATION NUMBER: US 60/292,336
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                                                                                                                        CTGAATGGGACTATCCACAGCACAAGTGAGGCAGACACAGAACCCTGTGTGGGATGGCTGG 360
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Johnson, Kor,
The Arthur
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THER INFORMATION: Clone ID: 08-LIB34-012-Q1-E1-B7
US-09-983-965-4911
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SEQ ID NO 4511
LENGTH: 370
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                                                                                                                                    Sequence 4, Application US/09798743A Patent No. US20020099093A1
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APPLICANT: Byatt, John C.
APPLICANT: Myattalagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: MUCCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 37-21(10297)C
CURRENT APPLICATION NUMBER: US/09/983,965
CURRENT FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: US 09/465,231
PRIOR FILING DATE: 1999-12-15
PRIOR FILING DATE: 1999-12-15
PRIOR FILING DATE: 1999-12-15
                                                                                                                GENERAL INFORMATION:
               APPLICANT: Nezu, Jun-ichi
APPLICANT: Ose, Asuka
TITLE OF INVENTION: SYSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
FILE REFERENCE: 06501-073001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/113,678 PRIOR FILING DATE: 1998-12-17
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CURRENT APPLICATION NUMBER: US/09/798,743A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          581
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57.6%;
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Pred. No. 2.9e-14;
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514 GAGTGGAACCTGGTGTGTGAGGACGACTGGAAGGCCCCACTCACAATCTCCTTGTTCTTC 573
                                                                                     460 ACTGGAATGCTGGTGGGAGGCATCATAGGTGGCCATGTCTCAGACAGGTGGCTGGTGGAA 519
                                                                                                                                                  574 graddraticitridacrectricarreadachacrercadachachragecadada 633
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                                                                                                                                                                                                                                                                                   634 AATGTGCTGTTCGTGACCATGGGCA 658
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 47, Application US/09764887
Patent No. US20020042096A1
GENERAL INFORMATION:
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Best Local Similarity 56.24
Matches 82, Conservative
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Best Local Similarity
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LENGTH: 796
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APPLICANT: Negal, Jun-ichi
APPLICANT: Ose, Asuka
TITLE OF INVENTION: SYSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
FILTE OF INVENTION: SYSTEMIC CARNITINE DEFICIENCY GENE AND USES THEREOF
CURRENT APPLICATION NUMBER: US/09/798,743A
CURRENT APPLICATION NUMBER: US/09/04853
PRIOR FILING DATE: 1999-09-07
PRIOR FILING DATE: 1999-09-07
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTIN VUM CURRENT OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA OF SEGULIA
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CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: PCT/JP99/04853
PRIOR FILING DATE: 1999-00-07
PRIOR FILING DATE: 1998-09-07
PRIOR FILING DATE: 1998-09-07
NUMBER OF SEQ ID NOS: 31
SEQ ID NO 4
LENGTH: 1888
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US-09-798-743A-4
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ORGANISM: Homo sapiens
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Matches 114; Conserv
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US-09-798-743A-2
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Pred. No. 0.0016;
0; Mismatches 64; Indels 0
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
TITLE REPERENCE: PAIN
CURRENT APPLICATION NUMBER: US/09/764,887
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 658
SOFTWARE: Patentin Ver. 2.0
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Patent No. US20020042096A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PAll3
CURRENT APPLICATION NUMBER: US/09/764,887
CURRENT PALLING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 658
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LA ROSE, Thomas J.

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(15401)B
CURRENT APPLICATION NUMBER: US/09/878,574
CURRENT FILING DATE: 2001-12-21
PRIOR APPLICATION NUMBER: 09/333,535
PRIOR FILING DATE: 1999-06-14
NUMBER OF SEQ ID NOS: 15775
SEQ ID NO 8151
LENGTH: 275
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APPLICANT: Clausen, Ib Groth
TITLE OF INVENTION: Methods For Monitoring Multiple Gene
TITLE OF INVENTION: Expression
FILE REFERENCE: 10085,500-US
CURRENT APPLICATION NUMBER: US/09/974,300
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: 09/680,598
PRIOR APPLICATION NUMBER: 09/680,598
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 8481
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US-09-974-300-945
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; NAME/KEY: misc feature
; LOCATION: (1)...(810)
; OTHER INFORMATION: n = A,T,C or G
US-09-974-300-945
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 4.8%;
Best Local Similarity 52.5%;
Matches 84; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     155 AGAGAAGAAGCGAAGGCAAATCAAAATGCATCATGGGCTA 194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      35 GAATCACCACGTTTTTTAATAAAGAACAATAAACTTGATGAAGCTCGCAAGGTGTTGAGC 94
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Search completed: January 6, 2003, 22:51:06 Job time : 43.0945 secs

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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/2/ina/FCTUS_COMB.seq:*

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Copyright (c) 1993 - 2003 Compugen Ltd.
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RESULT 2 US-09-330-245A-1

Sequence 1, Application US/09330245A; Patent No. 6432631; GENERAL INFORMATION:
APPLICANT: GILEAD SCIENCES, INC. et al.
TITLE OF INVENTION: NOVEL GENE ENCODING ORGANIC ANION TRANSPORTER FILE REFERENCE: 240.1PCnew
CURRENT APPLICATION NUMBER: US/09/330,245A
CURRENT FILING DATE: 1999-06-10

| 28 31 3.9 1835 1 US-08-55-55-3 Sequence 3, Appli 30 31 3.9 1835 2 US-08-931-932 Sequence 3, Appli 30 31 3.9 62804 4 US-09-801-39 Sequence 3, Appli 31 30.8 31 3553 4 US-08-931-939-17 Sequence 176, Appli 32 30.4 3.8 335 4 US-09-962-315-16 Sequence 176, Appli 34 30.4 3.8 335 2 US-08-931-926-116 Sequence 176, Appli 35 30.4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-546-238-3 Sequence 3, Appli 30 4 3.8 3390 2 US-09-266-315-57 Sequence 3, Appli 4 US-09-266-315-57 Sequence 3, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 4 US-09-266-315-57 Sequence 57, Appli 5 Sequence 57, Appli 5 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 6 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09-266-315-57 Sequence 57, Appli 7 US-09- | | | | | | |
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Sequence 1, Application US/09572147

Patent No. 6420544

Patent No. 6420544

Patent No. 6420544

Patent No. 6420544

APPLICANT: Lin Yue

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APPLICANT: Lin Yue

TITLE OF INVENTION: POLYNUCLEOTIDE AND POLYPEPTIDE SEQUENCES

TITLE OF INVENTION: SCREENING METHODS THEREOF

FILE REFERENCE: GP-70622

CURRENT APPLICATION NUMBER: US/09/572,147

PRIOR APPLICATION NUMBER: 60/134,879
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 CTGGTTTTTATTCTTCCCTCTCTCATGTTATTAATCCCTCATATACTGCTAGAACTTT 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          181 GGTAATGAAACTGGAATCCTCAGTGAAGATGCCCTCTTGAGAATCTCTATCCCACTAGAC 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         301 CTGAATGGGACTATCCACAGCACAAGTGAGGCAGACACAGAACCCTGTGTGGATGGCTGG 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        361 GTATATGATCAAGCTACTTCCCTTCGACCATTGTGACTAAGTGGGACCTGGTATGTGAT 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              421 TATCAGTCACTGAAATCAGTGGTTCAATTCCTACTTCTGACTGGAATGCTGGAAGGC 480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 ATGGCCTTTGAGGAGCTCTTGAGTCAAGTTGGAGGCCTTGGGAGATTTCAGATGCTTCAT 60
                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Unknown Organism:This information of OTHER INFORMATION: is not available.

NAME/KEY: CDS

LOCATION: (263)...(1912)
US-09-330-245A-1
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 10.2%; Score 81.8; DB 4; Length 2123; Best Local Similarity 51.7%; Pred. No. 1.5e-16; Matches 262; Conservative 0; Mismatches 212; Indels 33.
                       PRIOR FILING DATE: 1998-06-11
PRIOR APPLICATION NUMBER: 60/132,267
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.0
LENGTH: 2123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      481 ATCATAGGTGGCCATGTCTCAGACAGG 507
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     60/088,864
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Unknown
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                                                                                                                                                                                                                                                                                                                        391 AITGIGACTAAGIGGGACCTGGTATGTGATTATCAGTCACTGAAATCAGTGGTTCAATTC 450
                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                  451 CTACTTCTGACTGGAATGCTGGTGGGAGGCATCATAGGTGGCCATGTCTCAGACAGG 507
                                                                                                                                                                                                                                                                                                                                                                                                                                      10.1%; Score 81; DB 4; Length 1638; 66.1%; Pred. No. 2.4e-16; Live 0; Mismatches 60; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/647,397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08647397
GENERAL INFORMATION:
APPLICANT: Beler, David R.
APPLICANT: Brady, Kevin P.
ITILE OF INVENTION: OSTEOCLAST TRANSPORTER
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSER: Wolf, Greenfield & Sacks, P.C.
STREET: Boston
CITY: Boston
PRIOR FILING DATE: 1999-05-19
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSEQ for Windows Version 3.0
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NAME: Gates, Edward R.
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: B080
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-720-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 617-720-2441
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2102 base pairs
                                                                                                                                                                                         Best Local Similarity 66.19
Matches 117; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
ORGANISM: Mus musculus
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STRANDEDNESS: double
                                                                                             TYPE: DNA ORGANISM: MUS MUSCULUS
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CLASSIFICATION: 424
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ANTI-SENSE: NO
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                                                       SEQ ID NO 1
LENGTH: 1638
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LOCATION:
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US-08-647-397-1
                                                                                                                                                                           Query Match
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             US-08-232-463-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                         TELEX: 899149
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      523
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                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: EP 91
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 349 TGCATCTGCCAAACGCCAGTCTTCCCAATGACACCCAGGGGGCCACCGAGCCATGCTTGG 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                293 TTCTTCACCTGAATGGGACTATCCACAGCACAAGTGAGGCAGACACAGAACCCTGTGTGG 352
                                                        TOPOLOGY: line
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1800 Dia
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                         TELEFAX:
                                                                                                                                                                                                                                                          NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                           STRANDEDNESS:
                                                                                                                                                                                                          TELEPHONE:
                                                                                                                            LENGTH:
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                                                                                                                            7218 base pairs
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                                                                                                                                                                                             (703)683-4109
                                                                            linear
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                                                                                                                                                                                                              (703)836-9300
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                                                                                           single
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5.8%;
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                                                                                                                                                                                                                                              30472/114 IMMU
Score 46.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suite 500
 DB 1;
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 Length 7218;
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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 5670367
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14,
             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1484 TGTAATTACCTATCTATGCAAGTAGTTAAAGAGATAGAAGAATTTGGTACRRRRRRRRR 1425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: DOKNEN, APPLICANT: SCHEIFLIGER, F.
APPLICANT: FALKNER, F.G.
APPLICANT: FALKNER, F.G.
APPLICANT: FALKNER, F.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/232,463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           415 TGTGATTATCAGTCACTGAAATCAGTGGTTCAATTCCTACTTCTGACTGGAATGCTGGTG
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                475 GGAGGCATCATAGGTGGCCATGTCTCAGACAGGTGGCTGGTGGAATCTGCTCGGTGGTTG 534
 SEQUENCE
                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 100
                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 22313-0299
                                                                                           NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30
                                     TELEPHONE: (703)683-4109
                                                                                                                                                          FILING DATE:
                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                        APPLICATION NUMBER:
                                TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AATGATTGCTACACAAAAGTGACCAAA 801
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GAGCTGGATGCAGCACAGACCAAAACTACTGTGTGTGACTTGTTCCGCAACCCCAGTATG 714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGAATAAAGAATGCTGAAGAAACCCTGAACATAGAGGTTGTAAGATCCACCATGCAGGAG 654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4, Application US/08232463
5670367
CHARACTERISTICS:
                               899149
                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Foley & Lardner
1800 Diagonal Road,
                                                               (703)836-9300
                                                                                                                                                           26-AUG-1991
                                                                                                                                                                          EP 91 114 300.6
                                                                                                                                                                                                         US/07/935,313
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                                                                                               30472/114 IMMU
                                                                                                                                                                                                                                                                                                                                                                                                                                               Suite 500
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RESULT 7
US-09-719-083A-1/C

is Sequence 1, Application US/09719083A

j Patent No. 6451568

j GENERAL INFORMATION:
    APPLICANT: MCMaster, Christopher R
    APPLICANT: Anette, Henneberry L
    TITLE OF INVENTION: Choline/ethanolaminephosphotransferase: Synthesis of
    TITLE OF INVENTION: Phosphatidylcholine, Phosphatidylethanolamine, and
    TITLE OF INVENTION: Platelet Activating Factor
    TITLE OF INVENTION: Platelet Activating Factor
    TITLE OF INVENTION: Platelet Activating Factor
    FILE REFERENCE: 84372-203
    CURRENT FAPLICATION WUMBER: US/09/719, 083A
    PRIOR APPLICATION NUMBER: US60/088,379
    PRIOR FILING DATE: 1999-06-08
    PRIOR FILING DATE: 1999-06-08
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                                                                                                                                                                                                                                                                                                                                     81 TCTCATGTTATTAATCCCTCATATACTGCTAGAGAACTTTGCTGCAGCCATTCCTGGTCA 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  141 TCGTTGCTGGGTCCACATGCTGGACAATAATACTGGATCTGGTAATGAAACTGGAATCCT 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1852 TICCACCTCACAAGGTAGTGGGATGTATCTGTCACAAGCTGAATTCCTCAGTAAAGAA 1793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     201 CAGTGAAGATGCCCTCTTGAGAATCTCTATCCCACTAGACTCAAATCTGAGGCCAGAGAA 260
                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                  21 GAGTCAAGTTGGAGGCCTTGGGAGATTTCAGATGCTTCATCTGGTTTTTATTCTTCCCTC 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91 TTAATCCCTCATATACTGCTAGAAACTTTGCTGCAGCCATTCCTGGTCATCGTTGCTGG 150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
4.4%; Score 35.2; DB 4; Length 2051;
Best Local Similarity 53.7%; Pred. No. 0.22;
Matches 73; Conservative 0; Mismatches 63; Indels 0
                                                                                                                                                                                               ch 4.5%; Score 36; DB 1; Length 7218; l Similarity 6.4%; Pred. No. 0.24; l8; Conservative 147; Mismatches 117; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: transit_peptide
LOCATION: (129)..(1379)
CORPER INFORMATION: Open reading frame encoding hCEP1 peptide
US-09-719-083A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 261 GTGTCGTCGCTTTGTCCATCCCCAGTGGCAGCTTCTTCACCT 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: human Jurkat T cell cDNA library
7218 base pairs
                    ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; IMMEDIATE SOURCE:
; CLONE: PTZGPL-F1S
US-08-232-463-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO.
LENGTH: 2051
                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                 Matches
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Sequence 15, Application US/07906871
Patent No. 5340739
GENERAL INFORMATION:
APPLICANT: Stevens, Richard L.
APPLICANT: Avaham, Shalom
TILLE OF INVENTION: HEMATOPOIETIC CELL SPECIFIC
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATORY ELEMENTS OF SERGLYCIN AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSE: Sterne, Kessler, Goldstein & Fox
STREET: 1225 Connecticut Avenue, N.W., Suite 300
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/906,871
FILING DATE: 19920103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/816,289
FILING DATE: US JAN 1992
PRIOR APPLICATION NUMBER: US/07/635,544
FILING DATE: 18-JAN-1991
PRIOR APPLICATION NUMBER: PCT/US89/03051
FILING DATE: 13-JUL-1989
PRIOR APPLICATION NUMBER: PCT/US89/03051
FILING DATE: 13-JUL-1988
APPLICATION NUMBER: US/07/224,035
FILING DATE: 13-JUL-1988
ATTORNEY-AGENT INFORMATION:
NAME: CIMBAIA, Michele A
REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0627.2830004
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEBOOME. (2010)833-7533
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TELEPAX: (202)833-8716
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: NUCLEIC ACID
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16397..17327
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                        USA
                   US-07-906-871-15/c
                                                                                                                                                                                                                                                                                                                                                                                           20036
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/09081320 Patent No. 6093544
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14758 GGTCGTCGAGAGAAAGCACGTAGATAGAGAAGGGGGCCCAGGGCCAGAGCCAGGGTCACAC 14699
                                                                      Matches
                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14638 TAAGGCAGGGCATGGAAGCCTCAAGAAAAAA 14608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14698 CAACATGITAAATACTGAGCAGAGAAAAAAGCACAACCAGAGAAGTAGGAAGAACAGGA 14639
                                                                                                                                                                                                                                                                                      TELEFAX: (716) 263-1600 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Meng, Baozhong
TITLE OF INVENTION: RUDESTRIS STEM PITTING ASSOCIATED VIRUS
TITLE OF INVENTION: NUCLEIC ACIDS, PROTEINS, AND THEIR USES
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 587 GCACAAATGGAATAAAGAATGCTGAAGAAACCCCTGAACATAGAGGTTGTAAGATCCACCA 646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     527 GGTGGTTGATAATCACCAATAAACTAGATGAGGGCTTAAAGGCACTTAGAAAAGTTGCAC 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Gonsalves, Dennis
                                                                                                                                                                                                                                                                                                                                                        NAME: Goldman, Michael L.
REGISTRATION NUMBER: 30,727
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 6
FILING DATE: 20-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 17-DEC-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                        MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                y Match 4.3%; Score 34.2; Di
Local Similarity 51.7%; Fred. No. 1.5;
hes 78; Conservative 0; Mismatches
559 GGCTTAAAAGCACTTAGAAAAGTTGCACGCACAAATGGAATAAAGAATGCTGAAGAAACC 618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                 / Match 4.1%; Score 33.2; DB 3; Length 8743; Local Similarity 50.0%; Pred. No. 2.2; Local Similarity 50.0%; Pred. No. 2.2; Local Similarity 0; Mismatches 83; Indels 0
                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGCAGGAGGAGCTGGATGCAGCACCAGACCAA 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rochester
                                                                                                                                                                                                                          nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Nixon, Hargrave, Devans & Doyle LLP Clinton Square, P.O. Box 1051
                                                                                                                                                                                                                                                                    8743 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.A.
                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US 60/047,147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       us 60/069,902
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                                                                                                                                                                                                                                                                                                                                                                                                                       30,727
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US-09-707-780-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Gonsalves, Dennis
APPLICANT: Gonsalves, Dennis
APPLICANT: Meng, Baozhong
TITLE OF INVENTION: RUPESTRIS STEM PITTING ASSOCIATED VIRUS
TITLE OF INVENTION: NUCLEIC ACIDS, PROTEINS, AND THEIR USES
FIT DEFERRACE, 07678/035006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local Similarity 50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09707780 Patent No. 6399308
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CURRENT FILING DATE: 2000-05-18
PRIOR APPLICATION NUMBER: 60/047,147
PRIOR ETILING DATE: 1997-05-20
PRIOR APPLICATION NUMBER: 60/069,902
PRIOR APPLICATION NUMBER: 60/069,902
PRIOR PILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 09/081,320
PRIOR PILING DATE: 1998-05-19
PRIOR FILING DATE: 1998-05-19
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APPLICANT: Meng, Baozhong
TITLE OF INVENTION: RUPESTRIS STEM PITTING ASSOCIATED VIRUS
TITLE OF INVENTION: NUCLEIC ACIDS, PROTEINS, AND THEIR USES
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-12-17
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/707,780
CURRENT FILING DATE: 2000-11-07
PRIOR APPLICATION NUMBER: 09/081,320
PRIOR FILING DATE: 1998-05-19
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                                                                                                                                                 PRIOR FILING DATE: 1998-05-19
PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 07678/035006
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1904 AATTGGTAGACAGAAAAATATTATCTAAACCAGATACGTCTAAAAACAGAAATAGAGATTA 1963

1964 AAGAAGATTTAGTAAAACGAATTAATTGAAATACCAATAGTTTCGAAGATATAATGACTA 2023

668 CACAGACCAAAACTACTGTGTGTGACTTGTTCCGCAACCCCAGTATGCGTAAAAGGATCT 2024 TACCTACCAGTATCCCTATGAGATA-TTTTTAAACGGTACTAAGAGAAAAGTATCT

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608 CTGAAGAAACCCTGAACATAGAGGTTGTAAGATCCACCATGCAGGAGGAGCTGGATGCAG 667

2083 AAAGCIGIAGAITITITCCAGAAIGGAIATIAAGGGAGAIGAIATIAGCAAAAIGGGAAIA 2142

788 CAAAAGTGACCAAA 801

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728 GTATCCTGGTATTTTTGAGAAAAAATCTCAAGGAAAAGGCATAAAATGATTGCTACA 787

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                                                                                                                                                                                                                                                                                                                                                                      619 CTGAACATAGAGGTTGTAAGATCCACCATGCAGGAGGAGCTGGATGCAGCACAGACCAAA 678
                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                        ch 4.1%; Score 33.2; DB 4; Length 8743; 1 Similarity 50.0%; Pred. No. 2.2; 83; Conservative 0; Mismatches 83; Indels 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Paoletti, Enzo
APPLICANT: Pincus, Steven E.
APPLICANT: Cox, William I.
APPLICANT: Kauffman, Elizabeth K.
TITLE OF INVENTION: Recombinant Poxvirus - Cytomegalovirus,
TITLE OF INVENTION: Compositions and Uses
NUMBER OF SEQUENCES: 190
CORRESPONDER ADDRESS:
ADDRESSEE: Curries, Morris & Safford, P.C.
STREET: S30 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7884 CCTACTCTCAGTGGCATTCTTGCCAAACGCAAGAGGATTATAGAGA 7929
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MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: The PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/658,665
FILING DATE: 05-JUN-1996
CLASSIFICATION: 42-1996
                                                                        ; TYPE: DNA
; OGANISM: Rupestris stem pitting associated virus
VS-09-707-180-1
  FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 454310-2720.1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 178, Application US/08658665
Patent No. 5997878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Frommer Esg., William S. REGISTRATION NUMBER: 25,506
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INFORMATION FOR SEQ ID NO: 178:
SEQUENCE CHARACTERISTICS:
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US-08-658-665-178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 5798 base pairs
TYPE: nucleic acid
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                                                 LENGTH: 8743
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Best Local Simi
Matches 122;
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SOFTWARE:
                       SEQ ID NO 1
                                                                                                                                                                           Query Match
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1904 AATTGGTAGACAGAAAATATTATCTAAACCAGATACGTCTAAAACAGAAATAGAGATTA 1963
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                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: RESTENOSIS/ATHEROSCLEROSIS DIAGNOSIS, TITLE OF INVENTION: PROPHYLAXIS AND THERAPY NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/796,101
FILING DATE: 05-FEB-1997
ATTORNEY.ACCURATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                  E: CURTIS, MORRIS & SAFFORD, P.C.
530 FIFTH AVENUE
                                                                                                          Sequence 39, Application US/08796101
Patent No. 6183752
GENERAL INFORMATION:
APPLICANT: EPSTEIN, STEPHEN E.
APPLICANT: SPERIN, TOREN
APPLICANT: SPERIN, EDITH
APPLICANT: SPEIN, EDITH
APPLICANT: ZHOU, YI FU
APPLICANT: ZHOU, YI FU
APPLICANT: ENDILE, LORNE
APPLICANT: PINCUS, STEVEN
APPLICANT: PINCUS, STEVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: KOWALSKI, THOMAS J.
REGISTRATION NUMBER: 32,147
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 764-5574
INFORMATION FOR SEQ ID NO: 39:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: DNA (genomic) US-08-796-101-39
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nucleic acid
EDNESS: single
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2143 AAACACGGAGAAAA 2156
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                                                                                   US-08-796-101-39
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Mismatches 131; Indels

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US-07-923-095-1
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                                   Matches
                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Sokol Dr., Patricia TAPPLICANT: Ziai Dr., Mohammad R.
                                                                                                                                                                                                                                                                                                              TELEFAX: 203-321-2971
TELEX: 710-474-4059
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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APPLICATION NUMBER: US/07/923,095
FILING DATE: 19920730
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                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                HYPOTHETICAL: I
                                                                                                                                                                                                               MOLECULE TYPE: CL
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NAME: Lowney Dr., Karen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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343 CCCTGTGTGGATGGCTGGGTATATGATCAAAGCTACTTCCCTTCGACCATTGTGACTAAG 402
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                                                                                                                 NAME/KEY:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                               TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 06904-0060
                                                                                                                                                                   ORGANISM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                 1730 base pairs
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                                                 52.2%;
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                                                 Score 31.6;
Pred. No. 3;
                                  Mismatches
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                                                                Length 1730;
                                  Indels
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US-08-229-511-1
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                                                                                                                                                   Query Match
Best Local
                                                                                                                                   Matches
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                                                                1529
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1589
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                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 30-JUL-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/922,307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sokol, Patricia Tyson
APPLICANT: Chandra, Manik
TITLE OF INVENTION: Apamin Binding Protein/Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1589
                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                    MOLECULE TYPE: CDNA
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                               403
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                                                                                                  343 CCCTGTGTGGGATGGCTGGGTATATGATCAAAGCTACTTCCCTTCGACCATTGTGACTAAG 402
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                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Carroll, Alice O. REGISTRATION NUMBER: 33
                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                 NAME/KEY:
                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                   LOCATION:
                                                                                                                                                                                                                                                                                                                                   ENGTH:
TGGGAGATGCAAGATGAGTACCAGGGGGGTGGTTAGGAATAATTTCTAAGTTTTTCCACCT 1648
                             TGGGACCTGGTATGTGATTATCAGTCACTGAAATCAGTGGTTCAATTCCTACTTCTGACT 462
                                                                CCCTTTATGGTGGGCTTTGTTTTTTTAAAGCCACGGACAATGGCACAGCTTACCTCAG 1588
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                                                                                                                                                                                                                                                                                                                                  1730 base pairs
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                                                                                                                                   Conservative
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                                                                                                                                                 3.9%;
52.2%;
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                                                                                                                                                 Score 31.6;
Pred. No. 3;
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                                                                                                                               Mismatches
                                                                                                                                                                 DB 1; Length 1730;
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Search completed: January 6, 2003, 22:45:13 Job time : 60.3264 secs